## <213> Homo sapiens

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  Tyr Asn Gln Glu Glu Asn Thr Ser Ser Thr Leu Thr His Ala Glu Asn
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Asp Cys Tyr Val Gly Asp Gly Tyr Ser Tyr Arg Gly Lys Met Asn Arg
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Gln Glu Asn Tyr Asn Met Phe Met Glu Asp Ala Glu Thr His Gly Ile
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Glu	Leu	Pne	ALA	112	Arg	Pne	Leu	птэ	113	001	110			113	5
	<b>.</b>		*	Cln		Leu	G111	Ara			Glu	Asp	Ser		
Ser	Lys	Arg	114	GIII	nis	Leu	GIU	114	E Dys	O.L.	014		115		
T1 -	C 0 10	C1.	114	דום	Tur	Gly	Lvs			Ser	Glu	Gly	Ala	Asn	Ser
116	Ser	115		110	-1-	017	116					116	5		
mhu	The	113	Car	Tla	G1n	Glu	Pro	Val	Val	Leu	Phe	His	Ser	Arg	Phe
1111	117		001			117	5				118	0			
Mot	Glu	T.e.u	Thr	Ara	Met	Gln	Gln	Lvs	Lys	Lys	Glu	Lys	Asp	Gln	Lys
110	_				119	O				119	5				1200
Dro	Lvs	Glu	Val	Glu	Lvs	Gln	Glu	Asp	Thr	Glu	Asn	His	Pro	Lys	Thr
				120	5				121	0				121	5
Pro	Glu	Ser	Ala	Pro	Glu	Asn	Lys	Asp	Ser	Glu	Leu	Lys	Thr	Pro	Pro
			122	0				122	5				123	U	
Ser	Val	Gly	Pro	Pro	Ser	Val	Thr	Val	Val	Thr	Leu	Glu	Ser	Ala	Pro
		123	5				124	0				124	5		
Ser	Ala	Leu	Glu	Lys	Thr	Thr	Gly	Asp	Lys	Thr	Val	Glu	Ala	Pro	Leu
	125	0				125	5				126	0			
Val	Thr	Glu	Glu	Lys	Thr	Val	Glu	Pro	Ala	Thr	Val	Ser	Glu	Glu	Ala
126	5				127	0				127	5				1280
Lys	Pro	Ala	Ser	Glu	Pro	Ala	Pro	Ala	Pro	Val	Glu	Gln	Leu	Glu	Gln
				128	5				129	0				129	5
Val	Asp	Leu	Pro	Pro	Gly	Ala	Asp	Pro	Asp	Lys	Glu	Ala	Ala	Met	Met
			130	00				130	15				131	U	
Pro	Ala	Gly	Val	. Glu	ı Glu	ı Gly	Ser	Ser	: Gly	Asp	Gln	Pro	Pro	Tyr	Leu
							132	0				132	5		
		131	.5												
Asp		131 Lys	.5 Pro	Pro	Thr	Pro	Gly	Ala	Ser	Phe	Ser	Glr	Ala	Glu	Ser
	Ala	Lys	Pro			133	Gly 5	Ala			134	0			
	Ala	Lys	Pro		ı Pro	133 Asp	Gly 5	Ala		Pro	Leu Leu	0			Ala
Asr	Ala 133 Val	Lys 0 Asp	Pro	Glu	1 Pro	133 Asp	Gly 5 Ser	Ala Thr	Glr	135	134 Leu 55	o Ser	Lys	Pro	

	1365	1370		1375
Thr Ala Asp Ala (	Glu Pro Asp	Ala Asn Gln 1385	Lys Ala Glu	Ala Ala Pro 1390
Glu Ser Gln Pro	Dro Ala Cer		Glu Val Asp	Pro Pro Val
1395	FIO AIA DEL	1400	1405	
Ala Ala Lys Asp				
	1415		1420	270
1410 Val Gln Ala Ala	Ala Wal Cor	Tle Val Glu		Thr Arg Lvs
	1430	iie vai oiu	1435	1440
1425 Ser Glu Arg Ile 2	Nam New Clu			
	1445	1450	Arg Der Hon	1455
Gly Glu Ala Gln				
1460		1465	nee ora mra	1470
Thr Arg Thr Ala	Cam Iva Acn			
1475	ser bys ass.	1480	1485	
Pro Ser Leu Pro				
1490	1495		1500	
Tyr Ala Thr Met				Lvs Glu Pro
1505	1510		1515	1520
Val Glu Gln Pro				Glu Leu Gln
	1525	1530		1535
Glu Ala Ala Ala	Val Pro Thr			Pro Pro Lys
1540		1545		1550
Thr Arg Arg Arg	Ala Asp Glu	Glu Glu Glu	Asn Glu Ala	Lys Glu Pro
1555		1560	1565	, -
Ala Glu Thr Leu	Lvs Pro Pro	Glu Gly Trp	Arg Ser Pro	Arg Ser Gln
1570	1579	5	1580	
Lys Thr Ala Ala	Gly Gly Gly	Pro Gln Gly	Lys Lys Gly	Lys Asn Glu
1585	1590		1595	1600
Pro Lys Val Asp	Ala Thr Arg	Pro Glu Ala	Thr Thr Glu	Val Gly Pro
	1605	1610	)	1615
Gln Ile Gly Val	Lys Glu Ser	Ser Met Glu	Pro Lys Ala	Ala Glu Glu
1620	)	1625		1630
Glu Ala Gly Ser	Glu Gln Lys	Arg Asp Arg	Lys Asp Ala	Gly Thr Asp
1635		1640	1645	5
Lys Asn Pro Pro	Glu Thr Ala	Pro Val Glu	Val Val Glu	Lys Lys Pro
1650	165		1660	
Ala Pro Glu Lys	Asn Ser Lys	Ser Lys Arg	Gly Arg Ser	Arg Asn Ser
1665	1670		1675	1680
Arg Leu Ala Val				Asp Ala Ala
	1685	1690		1695
Val Ser Pro Arg	Gly Ala Ala		Gly Glu Arg	Glu Ser Gly
1700		1705		1710
Val Val Ala Val	Ser Pro Glu			
1715		1720	1725	
Gly Leu Ser Ser			Val Asp Pro	Asp Lys Glu
1730	173		1740	
Pro Glu Lys Glu		Ala Ser Gly	Pro Ser Pro	Glu Ala Thr
1745	1750		1755	1760
Gln Leu Ala Lys				His Ile Ala
	1765	177		1775
Lys Leu Ala Glu			Tyr Lys Ala	Asp Ala Pro
1780	0	1785		1790
Glu Gly Leu Ala	Pro Glu Asp	Arg Asp Lys	Pro ALA HIS	GIN ALA SET

1795																
1810																
Gly Glu Pro Glu Asn Phe Pro Ala Pro Pro Pro Tyr Pro Gly Glu Ser 1825	Glu	Thr	Glu	Leu	Ala	Ala	Ala	Ile	Gly	Ser	Ile			Asp	Ile	Ser
1825														_		
GIN Thr Asp Leu GIN Pro Pro Ala GIY Ala GIN Ala Leu GIN Pro Ser 1845  GIU GIU GIY Met GIU Thr Asp GIU Ala Val Ser GIY IL Leu GIU Thr 1860  GIU Ala Ala Thr GIU Ser Ser Arg Pro Pro Val Asn Ala Pro Asp Pro 1875  GIU Ala Ala Thr GIU Ser Ser Arg Pro Pro Val Asn Ala Pro Asp Pro 1875  Ser Ala GIY Pro Thr Asp Thr Lys GIU Ala Arg GIY Asn Ser Ser GIU 1895  Thr Ser His Ser Val Pro GIU Ala Lys GIY Ser Lys GIU Val GIU Val 1905  Thr Ser His Ser Val Pro GIU Ala Lys GIY Ser Lys GIU Val GIU Val 1905  Thr Leu Val Arg Lys Asp Lys GIY Arg GIN Lys Thr Thr Arg Ser Arg 1906  Thr Leu Val Arg Lys Asp Lys GIY Arg GIN Lys Thr Thr Arg Ser Arg 1940  Val Pro GIU Ser Asn GIN Ala GIN GIY Ser Pro Ala Ala Asn GIU 1955  GIY Thr Thr Val GIN His Pro GIU Ala Pro GIN GIU GIU Lys GIN Ser 1970  GIV Thr Thr Val GIN His Pro GIO Ala Ala Pro GIO GIU Lys GIN Ser 1970  GIV Thr Thr Val GIN His Pro GIO Ala Ala Pro GIN GIU GIU Lys GIN Ser 1970  GIV Thr Thr Val GIN His Pro GIN Ser Cys Thr Ser Asp Leu Ser 1990  Lys Ile Pro Ser Thr GIU Asn Ser Ser GIN GIU GIU Lys GIN Ser 1995  GIV Thr Thr Lys Ala Ser Val Pro Pro Asp Leu Pro Pro Pro Pro Pro 2020  GIN Pro Ala Pro Val Asp GIU GIU Pro GIN Ala Ag Phe Arg Val His 2035  Ser Ile Ile GIU Ser Asp Pro Val Thr Pro Pro Ser Asp Pro Ser Ile 2055  2055  2060  Val Ala Ser GIY GIY Hie Pro His GIN Ser Pro Pro Thr Lys Val Thr 2085  GIU Trp Ile Thr Arg GIN GIU GIU Pro Arg Ala GIN Ser Thr Pro Pro Ser Zilo  Pro Ala Leu Pro Pro Asp Thr Lys Ala Ser Asp Val Asp Thr Ser Ser Zilo  Pro Ala Leu Pro Pro Asp Thr Lys Ala Ser Asp Val Asp Thr Ser Ser Zilo  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ala Lys Leu Ser Pro Pro 2095  GIU Trp Ile Thr Arg GIN GIU GIU Pro Arg Ala GIN Ser Thr Pro Ser Zilo  Ser Val Thr Ser Thr Ser Val Thr Thr Ala GIU Pro Val Ser Zilo  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala GIU Pro Val Ser Zilo  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala GIU Pro Val Ser Zilo  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ala Ser Asp Val Asp Thr Ser Ser Zilo  Zilo Zilo Zilo Zilo Zilo Zilo Zilo Zilo			Pro	Glu				Ala	Pro				Pro	Gly	Glu	Ser
1845   1850   1850   1850   1850   1860	1825	5							_				_		_	
Glu Glu Gly Met Glu Thr Asp Glu Ala Val Ser Gly Ile Leu Glu Thr  1860 1865 1870  Glu Ala Ala Thr Glu Ser Ser Arg Pro Pro Val Asn Ala Pro Asp Pro  1875 1880 1985 1990  Ser Ala Gly Pro Thr Asp Thr Lys Glu Ala Arg Gly Asn Ser Ser Glu  1895 1990  Thr Ser His Ser Val Pro Glu Ala Lys Gly Ser Lys Glu Val Glu Val  1895 1990  Thr Ser His Ser Val Pro Glu Ala Lys Gly Ser Lys Glu Val Glu Val  1895 1990  Thr Ser His Ser Val Pro Glu Ala Lys Gly Ser Lys Glu Val Glu Val  1895 1990  Thr Leu Val Arg Lys Asp Lys Gly Arg Gln Lys Thr Thr Arg Ser Arg  1925 1930 1935  Arg Lys Arg Asn Thr Asn Lys Lys Val Val Ala Pro Val Glu Ser His  1940  Val Pro Glu Ser Asn Gln Ala Gln Gly Glu Ser Pro Ala Ala Asn Glu  1955 1960  Gly Thr Thr Val Gln His Pro Glu Ala Pro Gln Glu Glu Lys Gln Ser  1970 1975 1990  Glu Lys Pro His Ser Thr Pro Pro Gln Ser Gln Glu Glu Lys Gln Ser  1985 1990  Lys Ile Pro Ser Thr Glu Asn Ser Ser Gln Glu Ile Ser Val Glu Glu  2005 2020 2020  Gln Pro Ala Pro Val Asp Glu Glu Pro Gln Ala Arg Phe Arg Val His  2035 2040 2045  Ser Ile Ile Glu Ser Asp Pro Val Thr Pro Pro Ser Asp Pro Ser Ile  2050 2055 2060  Val Ala Ser Gly Gly Ile Pro Fis Gln Ser Fro Pro Thr Lys Val Thr  2065 2070 2075 2080  Glu Trp The Thr Leu Pro Ser Val Thr Ala Ala Lys Leu Ser Pro Pro  2065 2070 2075 2080  Glu Trp Ile Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser Lys  2016 1 Pro Thr Leu Pro Ser Val Thr Ala Ala Lys Leu Ser Pro Pro  2026 2075 2070 2075  Val Ala Ser Gly Gly Ile Pro Fis Gln Ser Fro Pro Thr Lys Val Thr  2085 2070 2075 2080  Glu Trp Ile Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser Ser  2125 2126  Fre Thr Leu Arg Lys Ile Leu Met Asp Pro Lys Tyr Val Ser Ala Thr  2130 2135 2140  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ala Glu Ser Pro Pro  2126 2127  Ser Thr Leu Arg Lys Ile Leu Met Asp Pro Pro Pro Pro Pro Pro Pro Val Asp Ser Lys  2169 2170 2175  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ala Glu Glu Pro Val Ser Ala Thr  2130 2135  Ser Val Thr Ser Thr Ser Val Thr Ala Pro Pro Pro Pro Pro Val Asp Ser Lys  2165 2170 2175  Ser Thr Leu Arg Lys Ile Leu Met	Gln	Thr	Asp	Leu			Pro	Ala				Ala	Leu			
1860							_						-1			
Glu Ala Ala Thr Glu Ser Ser Arg Pro Pro Val Asn Ala Pro Asp Pro 1875  Ser Ala Gly Pro Thr Asp Thr Lys Glu Ala Arg Gly Asn Ser Ser Glu 1890  Thr Ser His Ser Val Pro Glu Ala Lys Gly Ser Lys Glu Val Glu Val 1905  Thr Leu Val Arg Lys Asp Lys Gly Arg Gln Lys Thr Thr Arg Ser Arg 1920  Thr Leu Val Arg Lys Asp Lys Gly Arg Gln Lys Thr Thr Arg Ser Arg 1940  Val Pro Glu Ser Asn Gln Ala Gln Gly Glu Ser Pro Ala Ala Asn Glu 1955  Gly Thr Thr Val Gln His Pro Glu Ala Pro Gln Glu Glu Lys Gln Ser 1970  Glu Lys Pro His Ser Thr Pro Pro Gln Ser Cys Thr Ser Asp Leu Ser 1970  Glu Lys Pro His Ser Thr Glu Asn Ser Ser Gln Glu Glu Fro Pro Pro Pro Pro Pro Pro Pro Pro Pro P	Glu	Glu	Gly			Thr	Asp	GIu			Ser	GIY	шe	Leu	GIU	THE
1875		_				_	_						21-			Dwa
See Ala Gly Pro Thr Asp Thr Lys Glu Ala Arg Gly Asn Ser Ser Glu 1890	Glu	Ala			GIu	Ser				Pro	vaı	ASII			Asp	PIO
1890					mh	7.00				712	7.~~	Gly			cor	Glu
Thr Ser His Ser Val Pro Glu Ala Lys Gly Ser Lys Glu Val Glu Val 1910  Thr Leu Val Arg Lys Asp Lys Gly Arg Gln Lys Thr Thr Arg Ser Arg 1925  Arg Lys Arg Asn Thr Asn Lys Lys Val Val Ala Pro Val Glu Ser His 1940  Val Pro Glu Ser Asn Gln Ala Gln Glu Glu Ser Pro Ala Ala Asn Glu 1955  Gly Thr Thr Val Gln His Pro Glu Ala Pro Glu Glu Lys Gln Ser 1970  Glu Lys Pro His Ser Thr Pro Pro Gln Ser Cys Thr Ser Asp Leu Ser 1990  Lys Ile Pro Ser Thr Glu Asn Ser Ser Gln Glu Ile Ser Val Glu Glu Cys Pro Ala Ber Val Glu Glu Cys Pro Ala Ala Asn Glu 2005  Arg Thr Pro Thr Lys Ala Ser Val Pro Pro Asp Leu Pro	ser			Pro	IIII	Asp			GIU	ALG	ALG	1900	1	361	Jer	O_u
1905	Thr			Car	Val	Dro			Lve	Glv	Ser			Val	Glu	Val
Thr Leu Val Arg Lys Asp Lys Gly Arg Gln Lys Thr Thr Arg Ser Arg 1925 1935  Arg Lys Arg Asn Thr Asn Lys Lys Val Val Ala Pro Val Glu Ser His 1940 1945 1950  Val Pro Glu Ser Asn Gln Ala Gln Gly Glu Ser Pro Ala Ala Asn Glu 1955 1950  Gly Thr Thr Val Gln His Pro Glu Ala Pro Gln Glu Glu Lys Gln Ser 1970 1980  Glu Lys Pro His Ser Thr Pro Pro Gln Ser Cys Thr Ser Asp Leu Ser 1990 1995 2000  Lys Ile Pro Ser Thr Glu Asn Ser Ser Gln Glu Ille Ser Val Glu Glu Cys Gln Ser 2000  Arg Thr Pro Thr Lys Ala Ser Val Pro Pro Asp Leu Pro Pro Pro Pro Pro 2020  Gln Pro Ala Pro Val Asp Glu Glu Pro Gln Ala Arg Phe Arg Val His 2035 2055 2060  Ser Ile Ile Glu Ser Asp Pro Val Thr Pro Pro Ser Asp Pro Ser Ile 2055 2070 2075 2080  Val Ala Ser Gly Gly Ile Pro His Gln Ser Pro Pro Thr Lys Val Thr Pro Pro Pro Pro Pro 2055 2070 2075 2080  Glu Trp Ile Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser 2105 2105  Ser Val Thr Ser Thr Ser Val Thr Pro Pro Pro Pro Pro Ser Ser 2105 2105 2105  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ala Lys Leu Ser Thr Pro Ser 2115 2120 2125  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ala Glu Pro Val Ser Val Thr Pro Pro Pro Pro Pro Ser Ser Val Thr Pro Pro Pro Pro Pro Ser Ser Val Thr Pro Pro Val Asp Thr Ser Ser Val Thr Ser Thr Ser Val Thr Pro Pro Val Asp Thr Ser Ser Val Thr Pro Pro Val Asp Thr Ser Ser Val Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser 2105 2105  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser Ala Thr 2130 2155 2170  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Pro Pro Pro Val Asp Ser Lys 2165 2170 2175  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Val Thr Ash Ash Ser Glu 2195 2100  Pro Val Ile Ala Pro Vys Leu Val Ala Ala Asp Lys Glu Lys Val Ala 2195 2100  Pro Val Ile Ala Pro Vys Ile Thr Thr Ala Pro Pro Val Thr Ash Ash Ser Glu 2195  Pro Val Ile Ala Pro Vys Ile Thr Thr Val Ile Fro Val Thr Ash Ash Ser Cys 2165 2170  1216 Gln Ala Ser Glu Glu Lys Thr Ala Pro Pro Val Thr Ash Ash Ser Pro Val 2195 2105  Pro Val Ile Ala Pro Vys Ile Thr Thr Val Ile Val Ala 2195 2205  Pro Val Ile Ala Pro Vys Ile Thr Ser Val Il			1113	JCI	***				2,0							1920
1925   1930   1935   1936   1936   1936   1940   1945   1950   1940   1945   1950   1950   1950   1955   1955   1950   1955   1950   1955   1950   1955   1950   1955   1955   1950   1950   19			Val	Ara	Lvs			Glv	Ara	Gln	Lvs	Thr	Thr	Arq	Ser	Arq
1940					1929	5				1930	)				1935	5
1940	Arq	Lvs	Arg	Asn	Thr	Asn	Lys	Lys	Val	Val	Ala	Pro	Val	Glu	Ser	His
1955			~				•							1950	)	
Gly Thr Thr Val Gln His Pro Glu Ala Pro Gln Glu Glu Lys Gln Ser 1970  Glu Lys Pro His Ser Thr Pro Pro Gln Ser Cys Thr Ser Asp Leu Ser 1985  1990  Lys Ile Pro Ser Thr Glu Asn Ser Ser Gln Glu Glu Ile Ser Val Glu Glu Lys Ile Pro Ser Thr Glu Asn Ser Ser Gln Glu Ile Ser Val Glu Glu 2005  Arg Thr Pro Thr Lys Ala Ser Val Pro Pro Asp Leu Pro Pro Pro Pro Pro 2015  Arg Thr Pro Thr Lys Ala Ser Val Pro Pro Asp Leu Pro Pro Pro Pro Pro 2020  Gln Pro Ala Pro Val Asp Glu Glu Pro Gln Ala Arg Phe Arg Val His 2055  2056  Pro Ile Ile Glu Ser Asp Pro Val Thr Pro Pro Ser Asp Pro Ser Ile 2050  Pro Ile Pro Thr Leu Pro Ser Val Thr Ala Ala Lys Leu Ser Pro Pro Pro 2055  2070  Glu Trp Ile Thr Arg Gln Glu Glu Glu Fro Arg Ala Gln Ser Thr Pro Ser 2090  Glu Trp Ile Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser 2115  Ser Val Thr Ser Thr Ser Val Thr Lys Ala Ser Asp Val Asp Thr Ser Ser 2115  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser Ala Thr 2130  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser 2155  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser 2150  Ser Val Thr Ser Thr Ser Val Thr Ala Pro Pro Pro Pro Val Asp Ser Lys 2150  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Pro Pro Val Asp Ser Lys 2165  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Pro Val Thr Asn Asn Ser Glu 2195  Pro Val Ile Ala Pro Vyal Ile Thr Ser Val Ile Fra Arg Met Pro Val 2210  Pro Val Ile Ala Pro Vyal Ile Thr Fra Val Ile Ser Arg Met Pro Val 2210  Pro Val Ile Ala Pro Vyal Ile Thr Fra Val Ile Ser Arg Met Pro Val 2210	Val	Pro	Glu	Ser	Asn	Gln	Ala	Gln	Gly	Glu	Ser	Pro	Ala	Ala	Asn	Glu
1970																
Glu Lys Pro His Ser Thr Pro Pro Gln Ser Cys Thr Ser Asp Leu Ser 1995	Gly	Thr	Thr	Val	Gln	His	Pro	Glu	Ala	Pro	Gln			Lys	Gln	Ser
1995															_	_
Lys Tie Pro Ser Thr Glu Asn Ser Ser Gln Glu Ile Ser Val Glu Glu 2005  Arg Thr Pro Thr Lys Ala Ser Val Pro Pro Asp Leu Pro Pro Pro Pro Pro 2020  Gln Pro Ala Pro Val Asp Glu Glu Pro Gln Ala Arg Phe Arg Val His 2035  Ser Ile Ile Glu Ser Asp Pro Val Thr Pro Pro Ser Asp Pro Ser Ile 2050  Pro Tie Pro Thr Leu Pro Ser Val Thr Ala Ala Lys Leu Ser Pro Pro 2065  Pro Tie Pro Thr Leu Pro Ser Val Thr Ala Ala Lys Leu Ser Pro Pro 2065  Pro Tie Pro Thr Leu Pro Ser Val Thr Ala Ala Lys Leu Ser Pro Pro 2065  Val Ala Ser Gly Gly Ile Pro His Gln Ser Pro Pro Thr Lys Val Thr 2095  Glu Trp Ile Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser 2100  Pro Ala Leu Pro Pro Asp Thr Lys Ala Ser Asp Val Asp Thr Ser Ser 2110  Ser Thr Leu Arg Lys Ile Leu Met Asp Pro Lys Tyr Val Ser Ala Thr 2130  Ser Val Thr Ser Thr Ser Val Thr Thr Ala The Ala Glu Pro Val Ser 2145  Ser Val Thr Ser Thr Ser Val Thr Thr Ala The Ala Glu Pro Val Ser 2145  Ala Ala Pro Cys Leu His Glu Ala Pro Pro Pro Pro Val Asp Ser Lys 2160  11e Gln Ala Ser Glu Val Leu Val Ala Ala Asp Lys Glu Lys Val Ala 2195  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val Ala 2195  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2210  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2215  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2210			Pro	His				Pro	Gln				Ser	Asp	Leu	Ser
2005 2010 2015 2010 2015 2016 2016 2017 2016 2017 2017 2018 2018 2019 2019 2019 2019 2019 2019 2019 2019			_	_									0	17- 1	a1	
Arg Thr Pro Thr Lys Ala Ser Val Pro Pro Asp Leu Pro Pro Pro Pro 2020 2025 2030 2030 2035 2030 2030 2035 2030 2035 2030 2035 2035	Lys	Ile	Pro	Ser			Asn	ser	ser			TIE	ser	vai		
2020   2025   2030   2041   2041   2041   2041   2042   2045		m\	D	mla			Cor	17-1	Dro			Lan	Dro	Dro		
Gln Pro Ala Pro Val Asp Glu Glu Pro Gln Ala Arg Phe Arg Val His 2035  Ser Ile Ile Glu Ser Asp Pro Val Thr Pro Pro Ser Asp Pro Ser Ile 2055  Ser Ile Ile Glu Ser Asp Pro Val Thr Pro Pro Ser Asp Pro Ser Ile 2055  Ser Ile Pro Thr Leu Pro Ser Val Thr Ala Ala Lys Leu Ser Pro Pro 2055  Val Ala Ser Gly Gly Ile Pro His Gln Ser Pro Pro Thr Lys Val Thr 2035  Glu Trp Ile Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser 2010  Pro Ala Leu Pro Pro Asp Thr Lys Ala Ser Asp Val Asp Thr Ser Ser 2115  Ser Thr Leu Arg Lys Ile Leu Met Asp Pro Lys Tyr Val Ser Ala Thr 2135  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser 2155  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser 2155  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Pro Pro Val Asp Ser Lys 165  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Val Thr Asn Asn Ser Glu 2195  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Val Thr Asn Asn Ser Glu 2195  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Val Thr Asn Asn Ser Glu 2195  Lys Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2195  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 22105	Arg	THE	PIO			мта	ser				лър	Deu	FIU			110
2035 2040 2045  For Ile Ile Glu Ser Asp Pro Val Thr Pro Pro Ser Asp Pro	Gln	Dro	בות			Aen	Glu				Ala	Ara	Phe			His
Ser Ile Ile Glu Ser Asp Pro Val Thr Pro Pro Ser Asp Pro Ser Ile 2050 2055 2060 2060 2055 2060 2060 2055 2060 2060	GIII	FIU			•	p						3	2045	;		
2050 2055 2060  2070 2075 2070  2080 2075 2070  2080 2075 2070  2081 Ala Ser Gly Gly Ile Pro His Gln Ser Pro Pro Thr Lys Val Thr Ala Ala Lys Leu Ser Pro Pro 2080  Val Ala Ser Gly Gly Ile Pro His Gln Ser Pro Pro Thr Lys Val Thr Ala 1 ala Ser Ser Pro Pro Thr Lys Val Thr 2000  Glu Trp Ile Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser 2100  2100 2105 2105 2110  Pro Ala Leu Pro Pro Asp Thr Lys Ala Ser Asp Val Asp Thr Ser Ser 2115  2120 2125 2120  Ser Thr Leu Arg Lys Ile Leu Met Asp Pro Lys Tyr Val Ser Ala Thr 2130  2135 2150 2150  2150 2150 2150  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser 2145  2150 2155 2170  2155 2170  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Pro Val Asp Ser Lys 2165  Lys Pro Leu Glu Glu Ser Thr Ala Pro Pro Val Thr An Asn Ser Glu 2180  2180 2185 2180 2185  Pro Val Ile Ala Pro Val Ile Thr Ser Val Ile Ser Arg Met Pro Val 2195  Pro Val Ile Ala Pro Vs Ile Thr Ser Val Ile Ser Arg Met Pro Val 2215  Pro Val Ile Ala Pro Vs Ile Thr Ser Val Ile Ser Arg Met Pro Val 2215	Ser	Ile	Ile	Glu	Ser	Asp	Pro	Val	Thr	Pro	Pro	Ser	Asp	Pro	Ser	Ile
2055   2070   2075   2080						-										
Val Ala Ser Gly Gly Ile Pro His Gln Ser Pro Pro Thr Lys Val Thr 2095  Glu Trp Ile Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser 2100  Pro Ala Leu Pro Pro Asp Thr Lys Ala Ser Asp Val Asp Thr Ser Ser 2120  Ser Thr Leu Arg Lys Ile Leu Met Asp Pro Lys Tyr Val Ser Ala Thr 2130  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser 2145  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser 2145  Ala Ala Pro Cys Leu His Glu Ala Pro Pro Pro Pro Val Asp Ser Lys 2150  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Val Thr Asn Asn Ser Glu 2186  1le Gln Ala Ser Glu Val Leu Val Ala Ala Asp Lys Glu Lys Val Ala 2195  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Il Ser Arg Met Pro Val 2210  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Il Ser Arg Met Pro Val 2210  2215  2220  Pro Val Ile Ala Pro Lys 215 2220	Pro	Ile	Pro	Thr	Leu	Pro	Ser	Val	Thr	Ala	Ala	Lys	Leu	Ser	Pro	Pro
2095   2096   2095   2096   2097   2097   2098																
Glu Trp Ile Thr Arg Gln Glu Glu Pro Arg Ala Gln Ser Thr Pro Ser 2100 2105 2110	Val	Ala	Ser	Gly	Gly	Ile	Pro	His	Gln			Pro	Thr	Lys	Val	Thr
2100   2105   2110   2110   2125   2120   2125   2125   2125   2126   2126   2125   2126   2126   2126   2126   2126   2126   2126   2126   2126   2126   2126   2126   2126   2126   2126   2126   2126   2126   2125   2126   2125   2126   2125   2125   2126   2125   2125   2126   2125   2125   2126   2125   2125   2126   2125   2125   2126   2126   2125   2126																
Pro         Ala         Leu         Pro         Pro         Asp         Thr         Lys         Ala         Ser         Asp         Val         Asp         Thr         Ser         Ser           Ser         Thr         Leu         Met         Asp         Pro         Lys         Tyr         Val         Ser         Ala         Thr         Thr         Val         Thr         Thr         Ala         Ala         Ala         Glu         Pro         Val         Ser         Lys         Thr         Thr         Thr         Ala         Ala         Ala         Glu         Pro         Val         Asp         Ser         2150         2150         2155         2160         2160         Ala         Asp         Ser         Lys         2160         2150         2150         2175         2150         210         2150         210         210         2150         210	Glu	Trp	Ile			Gln	Glu	Glu			Ala	Gln	Ser			Ser
2115   2120   2125   2126   2125   2126   2135											_		_			
Ser Thr Leu Arg Lys Ile Leu Met Asp Pro Lys Tyr Val Ser Ala Thr 2130 2135 2140  Ser Val Thr Ser Thr Ser Val Thr Thr Ala Ile Ala Glu Pro Val Ser 2145 2150 2155 2160  Ala Ala Pro Cys Leu His Glu Ala Pro Pro Pro Val Asp Ser Lys 2165 2170 2175  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Val Thr Asn Asn Ser Glu 2180 2180  Ile Gln Ala Ser Glu Val Leu Val Ala Ala Asp Lys Glu Lys Val Ala 2195 2200 2205  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2210 2215 2220	Pro	Ala			Pro	Asp				Ser	Asp	Val	Asp	Thr	ser	ser
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$					_					_					21.	mla
Ser Val Thr Ser Thr         Ser Val Thr         Thr Ala         Ile Ala         Glu Pro         Val         Ser 2155         2155         2150         2150         2160         2160         2160         2160         2175	Ser			Arg	Lys	He			Asp	Pro	Lys			ser	ALA	THE
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$					mla sa	C			mh~	77-	т1 о			Dro	V=1	Ser
Ala Ala Pro Cys Leu His Glu Ala Pro Pro Pro Pro Val Asp Ser Lys 2165 2170 2175 Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Val Thr Asn Asn Ser Glu 2180 2185 2190 Ile Gln Ala Ser Glu Val Leu Val Ala Ala Asp Lys Glu Lys Val Ala 2195 2200 Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2210 2215 2220			mr	ser	Int			1111	IIII	мта			GIU	FIU	val	
2165 2170 2175  Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Val Thr Asn Asn Ser Glu 2180 2185 2190 2190  The Gln Ala Ser Glu Val Leu Val Ala Ala Asp Lys Glu Lys Val Ala 2195 2200 2205  Pro Val Ile Ala Pro Lys Thr Ser Val Ile Ser Arg Met Pro Val 2210 2215 2225 2226			Dwo	Crea	Lou			71-	Dro	Dro			Val	Asn	Ser	
Lys Pro Leu Glu Glu Lys Thr Ala Pro Pro Val Thr Asn Asn Ser Glu 2180 2195 2190  Ile Gln Ala Ser Glu Val Leu Val Ala Ala Asp Lys Glu Lys Val Ala 2195 2200 2205  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2210 2215 2220			PIO	Cys	neu			nia	FIU	FIO			•442	· · · · ·		-1-
2180 2185 2190  The Gin Ala Ser Glu Val Leu Val Ala Ala Asp Lys Glu Lys Val Ala 2195 2200 2205  Pro Val Ile Ala Pro Lys lle Thr Ser Val Ile Ser Arg Met Pro Val 2210 2215 2220			Len	Glu	Glu			Ala	Pro	Pro			Asn	Asn	Ser	Glu
Ile Gln Ala Ser Glu Val Leu Val Ala Ala Asp Lys Glu Lys Val Ala 2195 2200 2205  Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2210 2225 2225	nys	FIO	Deu			Lys	****	ALG			•					
2195 2200 2205 Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2210 2215 2220	Tla	Gln	Δla			Va 1	Len	Va l			Asp	Lvs	Glu	Lvs	Val	Ala
Pro Val Ile Ala Pro Lys Ile Thr Ser Val Ile Ser Arg Met Pro Val 2210 2215 2220	-10											-				
2210 2215 2220	Pro	Val			Pro	Lys	Ile			Val	Ile	Ser	Arg	Met	Pro	Val
Ser Ile Asp Leu Glu Asn Ser Gln Lys Ile Thr Leu Ala Lys Pro Ala																

										2225					2240
2225			_		2230	_				2235		<b>a</b> 1	T	17-1	
Pro	Gln	Thr	Leu			Leu	Val			Leu	Inr	сту	Leu	2255	ASII
				2245					2250			_			
Val	Ser	Leu	Val	Pro	Val	Asn	Ala	Leu	Lys	Gly	Pro	Val	Lys	GLY	Ser
			2260					2265					2270		
Val	Thr	Thr	Leu	Lys	Ser	Leu	Val	Ser	Thr	Pro	Ala	Gly	Pro	Val	Asn
		2275					2280					2285			
Val	T.011	Lare	Glv	Pro	Val	Asn	Val	Leu	Thr	Gly	Pro	Val	Asn	Val	Leu
val	2290					2295				2	2300	3			
					-1-			a1	The sec	Val			712	Dro	Gly
		Pro	var	ASII			vai	Gry	1111	2315		AIG	7.14	110	2320
2305					2310				_						
Thr	Val	Asn	Ala			Ser	Ala	Val	Asn	Ala	Thr	Ата	ser		
				2325					2330					2335	
Thr	Val	Thr	Ala	Gly	Ala	Val	Thr	Ala	Ala	Ser	Gly	Gly	Val	Thr	Ala
			2340	)				2345	5				2350	)	
Thr	Thr	Glv	Thr	Val	Thr	Met	Ala	Glv	Ala	Val	Ile	Ala	Pro	Ser	Thr
		2355					2360					2365			
T	Crra			7, 2007	7 l a				Glu	Asn	Ser			His	Pro
ьув			GIII	ALG		2379		A3**	u		2380	,			
	2370		_										a1	c	G1
		Met	Pro	Val			Asp	Arg	Pro	Ala	. ASP	Ата	GLY	Ser	GIY
2385	5				2390					2395					2400
Ala	Gly	Leu	Arg	Val	Asn	Thr	Ser	Glu	Gly	Val	Val	Leu	Leu	Ser	Tyr
				2405					2410					2419	
Ser	Gly	Gln	Lys	Thr	Glu	Gly	Pro	Gln	Arg	Ile	Ser	Ala	Lys	Ile	Ser
	2		2420			-		2425	5				2430	)	
Gln	Tle	Pro			Ser	Ala	Met	Asp	Ile	Glu	Phe	Gln	Gln	Ser	Val
GIII	110	243					2440					2445			
	T			17-1	Live	Dro			Val	Thr	Δla			Pro	Pro
ser			GIII	vai		2455		Ser	vai		2460		0211		
	2450	, .	_										.1-	mla -a	***
		Gly	Pro	GIn			Ala	GIA	Tyr	Ala	ASI	vai	ALA	Inr	nis
2469					2470					2475					2480
Ser	Thr	Leu	Val			Ala	Gln	Thr		Asn	Ala	Ser	Pro		
				248					2490					249	
Ser	Sar										T	Pro			
		Val	Lys	Ala	Asp	Arg	Pro	Ser	Leu	Glu	Lys		GIU	Pro	Ile
	501	Val	Lys 250	Ala	Asp	Arg	Pro	Ser 2509	Leu	Glu	ьys		2510	Pro	Ile
wie			250	0				2509	5				2510	)	
His		Ser	250 Val	0			Val	2509 Thr	5	Glu Gly		Thr	2510 Val	)	
	Leu	Ser 251	250 Val	Ser	Thr	Pro	Val 2520	2509 Thr	Gln	Gly	Gly	Thr 2525	2510 Val	Lys	Val
	Leu Thr	Ser 251 Gln	250 Val	Ser	Thr	Pro Thr	Val 2520 Pro	2509 Thr	Gln		Gly Val	Thr 2529 His	2510 Val	Lys	Val
Leu	Leu Thr 253	Ser 251: Gln	250 Val 5 Gly	Ser Ile	Thr Asn	Pro Thr 253	Val 2520 Pro	2509 Thr ) Pro	Gln Val	Gly Leu	Gly Val 254	Thr 2529 His	2510 Val S Asn	Lys Gln	Val Leu
Leu Val	Leu Thr 253 Leu	Ser 251: Gln	250 Val 5 Gly	Ser Ile	Thr Asn Ile	Pro Thr 2535 Val	Val 2520 Pro	2509 Thr ) Pro	Gln Val	Gly Leu Lys	Gly Val 2540 Lys	Thr 2529 His	2510 Val S Asn	Lys Gln	Val Leu Pro
Leu Val 254	Leu Thr 253 Leu	Ser 251: Gln Thr	250 Val 5 Gly Pro	Ser Ile Ser	Thr Asn Ile 255	Pro Thr 2539 Val	Val 2520 Pro Thr	2509 Thr Pro Thr	Gln Val Asn	Gly Leu Lys 2555	Gly Val 2540 Lys	Thr 2529 His Leu	2510 Val S Asn Ala	Lys Gln Asp	Val Leu Pro 2560
Leu Val 254	Leu Thr 253 Leu	Ser 251: Gln Thr	250 Val 5 Gly Pro	Ser Ile Ser	Thr Asn Ile 255	Pro Thr 2539 Val	Val 2520 Pro Thr	2509 Thr Pro Thr	Gln Val Asn	Gly Leu Lys	Gly Val 2540 Lys	Thr 2529 His Leu	2510 Val S Asn Ala	Lys Gln Asp	Val Leu Pro 2560
Leu Val 254	Leu Thr 253 Leu	Ser 251: Gln Thr	250 Val 5 Gly Pro	Ser Ile Ser	Thr Asn Ile 2556 Glu	Pro Thr 2539 Val	Val 2520 Pro Thr	2509 Thr Pro Thr	Gln Val Asn	Gly Leu Lys 2555 Gln	Gly Val 2540 Lys	Thr 2529 His Leu	2510 Val S Asn Ala	Lys Gln Asp	Val Leu Pro 2560 Gly
Leu Val 254 Val	Leu Thr 2530 Leu 5	Ser 251: Gln Thr	2500 Val 5 Gly Pro Lys	Ser Ile Ser Ile 256	Thr Asn Ile 2550 Glu	Pro Thr 253: Val O Thr	Val 2520 Pro Thr	Thr Pro Thr	Gln Val Asn Leu 2570	Gly Leu Lys 2555 Gln	Val 2540 Lys Pro	Thr 2529 His Leu Ala	2510 Val S Asn Ala Asn	Lys Gln Asp Leu 257	Val Leu Pro 2560 Gly
Leu Val 254 Val	Leu Thr 2530 Leu 5	Ser 251: Gln Thr	2500 Val 5 Gly Pro Lys Thr	Ser Ile Ser Ile 256:	Thr Asn Ile 2550 Glu	Pro Thr 253: Val O Thr	Val 2520 Pro Thr	2509 Thr Pro Thr Val	Gln Val Asn Leu 2570	Gly Leu Lys 2555 Gln	Val 2540 Lys Pro	Thr 2529 His Leu Ala	2510 Val Asn Ala Asn Lys	Lys Gln Asp Leu 257	Val Leu Pro 2560 Gly
Leu Val 254: Val Ser	Leu Thr 2530 Leu Thr Thr	Ser 251: Gln Thr Leu	2500 Val 5 Gly Pro Lys Thr 258	Ser Ile Ser Ile 2569 Pro	Thr Asn Ile 2556 Glu His	Pro Thr 253: Val Thr His	Val 2520 Pro Thr Lys	Thr Pro Thr Val Pro 258	Gln Val Asn Leu 2570 Ala	Gly Leu Lys 2559 Gln Leu	Val 2540 Lys Pro	Thr 2529 His Leu Ala Ser	2510 Val Asn Ala Asn Lys 2590	Lys Gln Asp Leu 257	Val Leu Pro 2560 Gly 5 Pro
Leu Val 254: Val Ser	Leu Thr 2530 Leu Thr Thr	Ser 251: Gln Thr Leu Leu	2500 Val 5 Gly Pro Lys Thr 258 Asn	Ser Ile Ser Ile 2569 Pro	Thr Asn Ile 2556 Glu His	Pro Thr 253: Val Thr His	Val 2520 Pro Thr Lys Pro	Thr Pro Thr Val Pro 258: Gly	Gln Val Asn Leu 2570 Ala	Gly Leu Lys 2555 Gln	Val 2540 Lys Pro	Thr 2525 His Leu Ala Ser	2510 Val Asn Ala Asn Lys 2590 Ala	Lys Gln Asp Leu 257	Val Leu Pro 2560 Gly 5 Pro
Leu Val 254: Val Ser Thr	Leu Thr 2533 Leu Thr Thr	Ser 251: Gln Thr Leu Leu Val 259	2500 Val 5 Gly Pro Lys Thr 258 Asn	Ser Ile Ser Ile 256: Pro His	Thr Asn Ile 2550 Glu His Val	Pro Thr 253! Val Thr His	Val 2520 Pro Thr Lys Pro Ser 2600	Thr Pro Thr Val Pro 2589 Gly	Gln Val Asn Leu 2570 Ala Pro	Leu Lys 2555 Gln Leu Ser	Val 2540 Lys Pro Pro	Thr 2525 His Leu Ala Ser Pro 2605	2510 Val Asn Ala Asn Lys 2590 Ala	Leu 257! Leu Asp Leu 257! Leu	Val Leu Pro 2560 Gly 5 Pro
Leu Val 254: Val Ser Thr	Leu Thr 2533 Leu Thr Thr	Ser 251: Gln Thr Leu Leu Val 259	2500 Val Gly Pro Lys Thr 258 Asn	Ser Ile Ser Ile 256: Pro His	Thr Asn Ile 2550 Glu His Val	Pro Thr 253: Val Thr His Pro	Val 2520 Pro Thr Lys Pro Ser 2600 Ala	Thr Pro Thr Val Pro 2589 Gly	Gln Val Asn Leu 2570 Ala Pro	Gly Leu Lys 2559 Gln Leu	Gly Val 2540 Lys Pro Pro Ile Ala	Thr 2523 His Leu Ala Ser Pro 2603 His	2510 Val Asn Ala Asn Lys 2590 Ala	Leu 257! Leu Asp Leu 257! Leu	Val Leu Pro 2560 Gly 5 Pro
Leu Val 254: Val Ser Thr	Thr 2530 Leu 5 Thr Thr Glu Val 261	Ser 251: Gln Thr Leu Leu Val 259: Ser	2500 Val 5 Gly Pro Lys Thr 258 Asn 5	Ser Ile Ser Ile 256: Pro His	Thr Asn Ile 2556 Glu His Val	Thr 253: Val Thr His Pro	Val 2520 Pro Thr Lys Pro Ser 2600 Ala	Pro Thr Val Pro 2589 Gly Lys	Gln Val Asn Leu 2570 Ala Fro Leu	Leu Lys 2555 Gln Leu Ser	Val 2540 Lys Pro Pro Ile Ala 2620	Thr 252: His Leu Ala Ser Pro 260: His	2510 Val Asn Ala Asn Lys 2590 Ala Ser	Lys Gln Asp Leu 257: Leu Asp Asp	Val Leu Pro 2560 Gly 5 Pro Arg
Leu Val 254: Val Ser Thr	Thr 2530 Leu 5 Thr Thr Glu Val 261	Ser 251: Gln Thr Leu Leu Val 259: Ser	2500 Val 5 Gly Pro Lys Thr 258 Asn 5	Ser Ile Ser Ile 256: Pro His	Thr Asn Ile 2556 Glu His Val	Thr 253: Val Thr His Pro	Val 2520 Pro Thr Lys Pro Ser 2600 Ala	Pro Thr Val Pro 2589 Gly Lys	Gln Val Asn Leu 2570 Ala Fro Leu	Leu Lys 2555 Gln Leu Ser	Val 2540 Lys Pro Pro Ile Ala 2620	Thr 252: His Leu Ala Ser Pro 260: His	2510 Val Asn Ala Asn Lys 2590 Ala Ser	Lys Gln Asp Leu 257: Leu Asp Asp	Val Leu Pro 2560 Gly 5 Pro Arg
Leu Val 254: Val Ser Thr	Thr 2530 Leu 5 Thr Thr Glu Val 2610 Ser	Ser 251: Gln Thr Leu Leu Val 259: Ser	2500 Val 5 Gly Pro Lys Thr 258 Asn 5	Ser Ile Ser Ile 256: Pro His	Thr Asn Ile 2556 Glu His Val	Thr 253: Val Thr His Pro Ala 261: Ser	Val 2520 Pro Thr Lys Pro Ser 2600 Ala	Pro Thr Val Pro 2589 Gly Lys	Gln Val Asn Leu 2570 Ala Fro Leu	Leu Lys 2555 Gln Leu Ser	Val 2540 Lys Pro Pro Ile Ala 2620 Ala	Thr 252: His Leu Ala Ser Pro 260: His	2510 Val Asn Ala Asn Lys 2590 Ala Ser	Lys Gln Asp Leu 257: Leu Asp Asp	Val Leu Pro 2560 Gly 5 Pro Arg
Leu Val 254: Val Ser Thr Thr	Leu Thr 2530 Leu Thr Thr Glu Val 261 Ser	Ser 251: Gln Thr Leu Val 259: Ser	2500 Val 5 Gly Pro Lys Thr 258 Asn 5 His	Ser Ile Ser Ile 256: Pro His Leu	Thr Asn Ile 2556 Glu His Val Ala Pro 263	Thr 253: Val Thr His Pro Ala 261: Ser	Val 2520 Pro Thr Lys Pro Ser 2600 Ala Ser	Thr Val Pro 258: Gly Lys	Gln Val Asn Leu 2570 Ala Pro Leu Pro	Leu Lys 2555 Gln Leu Ser Asp Arg 263	Val 2540 Lys Pro Pro Ile Ala 2620 Ala	Thr 2529 His Leu Ala Ser Pro 2609 His O Ser	2510 Val Asn Ala Asn Lys 2590 Ala Ser	Leu 2579 Leu Asp Asp Pro	Val Leu Pro 2560 Gly 5 Pro Arg Arg
Leu Val 254: Val Ser Thr Thr	Leu Thr 2530 Leu Thr Thr Glu Val 261 Ser	Ser 251: Gln Thr Leu Val 259: Ser	2500 Val 5 Gly Pro Lys Thr 258 Asn 5 His	Ser Ile Ser Ile 2566 Pro His Leu Gly	Thr Asn Ile 2555 Glu 5 His Val Ala Pro 263 Ala	Thr 253: Val Thr His Pro Ala 261: Ser	Val 2520 Pro Thr Lys Pro Ser 2600 Ala Ser	Thr Val Pro 258: Gly Lys	Gln Val Asn Leu 2570 Ala Pro Leu Pro Asn	Leu Lys 2555 Gln Leu Ser Asp Arg 2635 Ala	Val 2540 Lys Pro Pro Ile Ala 2620 Ala	Thr 2529 His Leu Ala Ser Pro 2609 His O Ser	2510 Val Asn Ala Asn Lys 2590 Ala Ser	Leu 2579 Leu Asp Asp Pro	Val Leu Pro 2560 Gly Fro Arg Arg Ser 2640 Ala
Val 254: Val Ser Thr Thr Pro 262 Ser	Thr 2533 Leu 5 Thr Glu Val 261 Ser 5 Thr	Ser 2511 Gln Thr Leu Val 259 Ser 0 Gly Ala	2500 Val 5 Gly Pro Lys Thr 258 Asn 5 His	Ser Ile Ser Ile 256: Pro His Leu Gly Thr 2644	Thr Asn Ile 2556 Glu 5 His Val Ala Pro 2633 Ala 5	Pro Thr 2533 Val Thr His Pro Ala 261: Ser	Val 2520 Pro 5 Thr Lys Pro Ser 2600 Ala 5 Ser	2509 Thr Pro Thr Val Pro 2589 Gly Lys Phe	Gln Val Asn Leu 2570 Ala Fro Leu Pro Asn 2650	Leu Lys 2555 Gln Leu Ser Asp Arg 2635 Ala	Val 2544 Lys Pro Pro Ile Ala 262 Ala 5	Thr 2529 His Dieu Ala Ser Pro 2600 His Dieu Val	2510 Val Asn Ala Asn Lys 2590 Ala Ser His	Leu 257! Leu Asp Pro Leu 265	Val Leu Pro 2560 Gly 5 Pro Arg Arg Ser 2640 Ala

2660		2665		2670
Ser Val Ile Met 2675	Pro Pro His	Ser Ile Thr 2680	Gln Thr Val 268	
His Leu Ser Gln	Gly Glu Val			
2690	269		2700	204 120 001
Ile Thr Tyr Ser				Arg Ala Pro
2705	2710	014 1114 200	2715	2720
Leu Gln Pro Gln		Val Ara Ala		
Leu GIN FIO GIN	2725	273		2735
Pro Gln Pro Ala				
2740		2745	Leu Ala Sel	2750
Pro Glu Glu Glu			Wal Ala Arm	
2755.	var mis iyr	2760	276	
Pro Val Gln Ser	Clu Val Leu			
2770	277		2780	Arg Lea His
Pro Tyr Thr Val				Dro Wie Wal
2785	2790	var Arg IIe	2795	2800
Thr Ala Val Ser		Ara Ala Ala		
IIII MIA VAI SEI	2805	281		2815
Pro Pro Ala Ser				
2820		2825	GIY LYS GIU	2830
			The Dro Ma	
Thr Pro Asp Ala 2835	DAR WIS WIS	2840	284	
Val Pro Val Pro	Ton Dre 31a			
2850	285		2860	GIY GIG AIG
Arg Ile Leu Thr				Lou Dro Lou
2865	2870	SEL ASII GIL	2875	2880
			2075	
	Wal Wal Thr	wie Cly Val	Cln Tle Val	
			Gln Ile Val	
	2885	289	0	2895
Gly Glu Leu Phe	2885 Gln Glu Tyr	289 Arg Tyr Gly	0	2895 Thr Tyr His
Gly Glu Leu Phe 2900	2885 Gln Glu Tyr )	289 Arg Tyr Gly 2905	0 Asp Ile Arg	2895 Thr Tyr His 2910
Gly Glu Leu Phe 2900 Pro Pro Ala Gln	2885 Gln Glu Tyr )	289 Arg Tyr Gly 2905 Thr Gln Phe	0 Asp Ile Arg Pro Ala Ala	2895 Thr Tyr His 2910 Ser Ser Val
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915	2885 Gln Glu Tyr ) Leu Thr His	Arg Tyr Gly 2905 Thr Gln Phe 2920	O Asp Ile Arg Pro Ala Ala 292	2895 Thr Tyr His 2910 Ser Ser Val
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser	2885 Gln Glu Tyr ) Leu Thr His Arg Thr Lys	Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro	2895 Thr Tyr His 2910 Ser Ser Val
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293	Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro	Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2950	Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 Gln Pro Val	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala 2960
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2950 Pro Cys Pro	Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 Gln Pro Val	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2950 Pro Cys Pro	Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 Gln Pro Val Pro Ser Gln 297	0 Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln 0	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln 2975
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser	2885 Gln Glu Tyr Leu Thr His 293 Gln Pro Pro 2950 Pro Cys Pro 2965 Lys Met Pro	Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 Gln Pro Val Pro Ser Gln 297 Gln Val Ser	0 Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln 0	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln 2975 Lys Gly Thr
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2980	2885 Gln Glu Tyr ) Leu Thr His 293 Gln Pro Pro 2950 Pro Cys Pro 2965 Lys Met Pro	Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 Gln Pro Val Pro Ser Gln 297 Gln Val Ser 2985	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln O Gln Glu Ala	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln 2975 Lys Gly Thr 2990
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2980 Gln Thr Gly Val	2885 Gln Glu Tyr ) Leu Thr His 293 Gln Pro Pro 2950 Pro Cys Pro 2965 Lys Met Pro	289 Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 Gln Pro Val Pro Ser Gln 297 Gln Val Ser 2985 Arg Leu Pro	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln O Gln Glu Ala Ala Gly Pro	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln 2975 Lys Gly Thr 2990 Ala Asn Arg
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2986 Gln Trd Gly Val 2995	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2950 Pro Cys Pro 2965 Lys Met Pro Glu Gln Pro	289 Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 Gln Pro Val Pro Ser Gln 297 Gln Val Ser 2985 Arg Leu Pro 3000	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln O Gln Glu Ala Ala Gly Pro 300	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala 2975 Pro Gly Gln 2975 Lys Gly Thr 2990 Ala Asn Arg 5
Gly Glu Leu Phe 2900 Pro Pro Ala Gln Pro Pro Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2980 Gln Thr Gly Val 2995 Pro Pro Glu Pro	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2955 Pro Cys Pro 2965 Lys Met Pro Glu Gln Pro His Thr Gln	289 Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 Gln Pro Val Pro Ser Gln 297 Gln Val Ser 2985 Arg Leu Pro 3000	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln O Gln Glu Ala Ala Gly Pro 3000 Ala Gln Ala	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala 2975 Pro Gly Gln 2975 Lys Gly Thr 2990 Ala Asn Arg 5
Gly Glu Leu Phe 2900 Pro Pro Ala Glin 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2986 Gln Thr Gly Val 2995 Pro Pro Glu Pro 3010	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2950 Pro Cys Pro 2965 Lys Met Pro Glu Gln Pro His Thr Gln 301	289 Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala5 Gln Pro Val Pro Ser Gln 297 Gln Val Ser 2985 Arg Leu Pro 3000 Val Gln Arg 5	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln O Gln Glu Ala Ala Gly Pro 300 Ala Gln Ala 3020	2895 Thr Tyr His 2910 Ser Ser Val 5 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln Gln 2975 Lys Gly Thr 2990 Ala Asn Arg 5 Glu Thr Gly
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2966 Gln Tro Gly Val 2995 Pro Pro Glu Pro 3010 Pro Thr Ser Phe	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2955 Lys Met Pro Glu Gln Pro His Thr Gln 301 Pro Ser Pro	289 Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala5 Gln Pro Val Pro Ser Gln 297 Gln Val Ser 2985 Arg Leu Pro 3000 Val Gln Arg 5	0 Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln O Gln Glu Ala Ala Gly Pro 300 Ala Gln Ala 3020 Ser Met Lys	2895 Thr Tyr His 2910 Ser Ser Val 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln 2975 Lys Gly Thr 2990 Ala Asn Arg 5 Glu Thr Gly Pro Asp Leu
Gly Glu Leu Phe 2900 Pro Pro Ala Glin 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2986 Gln Thr Gly Val 2995 Pro Pro Glu Pro 3010 Pro Thr Ser Phe 3025	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2955 Pro Cys Pro 2965 Glu Gln Pro His Thr Gln 301 Pro Ser Pro 3030	285 Arg Tyr Gln Phe 2905 Thr Gln Phe 2920 Thr Ala Ala 5 5 Gln Pro Val Pro Ser Gln 2985 Arg Leu Pro 3000 Val Gln Arg 5 Val Ser Val	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln Gln Glu Ala Ala Gly Pro 300 Ala Gln Ala 3020 Ser Met Lys 3035	2895 Thr Tyr His 2910 Ser Ser Val 5 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln 2975 Lys Gly Thr 2990 Ala Asn Arg 5 Glu Thr Gly Pro Asp Leu 3040
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2966 Gln Tro Gly Val 2995 Pro Pro Glu Pro 3010 Pro Thr Ser Phe	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2950 Pro Cys Pro 2965 Lys Met Pro Old Gln Pro His Thr Gln Pro Ser Pro 3030 Pro Thr Gln Fro	289 Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 Gln Pro Val Pro Ser Gln 2985 Arg Leu Pro 3000 Val Gln Arg 5 Val Ser Val Thr Ala Pro	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln O Gln Glu Ala Ala Gly Pro 300 Ala Gln Ala 3020 Ser Met Lys 3035 Lys Gln Pro	2895 Thr Tyr His 2910 Ser Ser Val 5 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln 2975 Lys Gly Thr 2990 Ala Asn Arg 5 Glu Thr Gly Pro Asp Leu 3040 Leu Phe Val
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2995 Gln Thr Gly Val 2995 Pro Pro Glu Pro 3010 Pro Thr Ser Phe 3025 Pro Val Ser Leu	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2955 Pro Cys Pro 2965 Glu Gln Pro Glu Gln Pro His Thr Gln 301 Pro Ser Pro 3030 Pro Thr Gln 3045	285 Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 55 5 Gln Pro Val Pro Ser Gln 297 Gln Val Ser 2985 Arg Leu Pro 3000 Val Gln Arg 5 Val Ser Val Thr Ala Pro 305	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln Gln Glu Ala Ala Gly Pro 300 Ala Gln Ala 3020 Ser Met Lys 3035 Lys Gln Pro 0	2895 Thr Tyr His 2910 Ser Ser Val 55 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln 2975 Lys Gly Thr 2990 Glu Thr Gly Pro Asp Leu 3040 Leu Phe Val 3055
Gly Glu Leu Phe 2900 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 2986 Gln Thr Gly Val 2995 Pro Pro Glu Pro 3010 Pro Thr Ser Phe 3025 Pro Val Ser Leu Pro Thr Thr Ser	2885 Gln Glu Tyr Leu Thr His Arg Thr Lys 293 Gln Pro Pro 2955 Pro Cys Pro 2965 Glu Gln Pro His Thr Gln 301 Pro Ser Pro 3030 Pro Thr Gln 3045	285 Arg Tyr Gly 2905 Thr Gln Phe 2920 Thr Ala Ala 5 5 Gln Pro Val Pro Ser Gln 2985 Gln Val Ser 2985 Arg Leu Pro 3000 Val Gln Arg 5 Val Ser Val Thr Ala Pro 305 Thr Pro Pro	O Asp Ile Arg Pro Ala Ala 292 Gln Gly Pro 2940 Gln Ser Thr 2955 Leu Gly Gln Gln Glu Ala Ala Gly Pro 300 Ala Gln Ala 3020 Ser Met Lys 3035 Lys Gln Pro 0	2895 Thr Tyr His 2910 Ser Ser Val 5 5 Pro Pro Glu Gln Pro Ala 2960 Pro Gly Gln Gln 2975 Lys Gly Thr 2990 Slus Ala Asn Arg 5 Glu Thr Gly Pro Ala Asn Arg 5 Glu Thr Gly Leu Phe Val 3055 Leu Pro His
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Val Ser Ser Leu Leu Gln Glu Glu Glu Pro Leu Ala Gly Gly Lys
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Pro Gly Ala Asp Gly Gly Ser Leu Glu Ala Val Arg Leu Gly Pro Ser
                        55
Ser Gly Leu Leu Val Asp Trp Leu Glu Met Leu Asp Pro Glu Val Val
                                        75
                    70
Ser Ser Cys Pro Asp Leu Gln Leu Arg Leu Leu Phe Ser Arg Arg Lys
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Gly Lys Gly Gln Ala Gln Val Pro Ser Phe Arg Pro Tyr Leu Leu Thr
            100
                               105
Leu Phe Thr His Gln Ser Ser Trp Pro Thr Leu His Gln Cys Ile Arg
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Val Leu Leu Gly Lys Ser Arg Glu Gln Arg Phe Asp Pro Ser Ala Ser
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                                            140
Leu Asp Phe Leu Trp Ala Cys Ile His Val Pro Arg Ile Trp Gln Gly
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                                       155
Arg Asp Gln Arg Thr Pro Gln Lys Arg Arg Glu Glu Leu Val Leu Arg
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Val Gln Gly Pro Glu Leu Ile Ser Leu Val Glu Leu Ile Leu Ala Glu
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Ala Glu Thr Arg Ser Gln Asp Gly Asp Thr Ala Ala Cys Ser Leu Ile
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                                                205
Gln Ala Arg Leu Pro Leu Leu Leu Ser Cys Cys Cys Gly Asp Asp Glu
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                                            220
Ser Val Arg Lys Val Thr Glu His Leu Ser Gly Cys Ile Gln Gln Trp
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                                        235
Gly Asp Ser Val Leu Gly Arg Arg Cys Arg Asp Leu Leu Gln Leu
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                                    250
Tyr Leu Gln Arg Pro Glu Leu Arg Val Pro Val Pro Glu Val Leu Leu
                                265
His Ser Glu Gly Ala Ala Ser Ser Ser Val Cys Lys Leu Asp Gly Leu
                                                285
                            280
Ile His Arg Phe Ile Thr Leu Leu Ala Asp Thr Ser Asp Ser Arg Ala
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                                           300
Leu Glu Asn Arg Gly Ala Asp Ala Ser Met Ala Cys Arg Lys Leu Ala
                    310
                                        315
Val Ala His Pro Leu Leu Leu Leu Arg His Leu Pro Met Ile Ala Ala
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Leu Leu His Gly Arg Thr His Leu Asn Phe Gln Glu Phe Arg Gln Gln
                                345
Asn His Leu Ser Cys Phe Leu His Val Leu Gly Leu Leu Glu Leu Leu
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Gln Pro His Val Phe Arg Ser Glu His Gln Gly Ala Leu Trp Asp Cys
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                                            380
Leu Leu Ser Phe Ile Arg Leu Leu Leu Asn Tyr Arg Lys Ser Ser Arg
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395
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385
His Leu Ala Ala Phe Ile Asn Lys Phe Val Gln Phe Ile His Lys Tyr
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Ile Thr Tyr Asn Ala Pro Ala Ala Ile Ser Phe Leu Gln Lys His Ala
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Asp Pro Leu His Asp Leu Ser Phe Asp Asn Ser Asp Leu Val Met Leu
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Lys Ser Leu Leu Ala Gly Leu Ser Leu Pro Ser Arg Asp Asp Arg Thr
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Asp Arg Gly Leu Asp Glu Glu Gly Glu Glu Glu Ser Ser Ala Gly Ser
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                    470
Leu Pro Leu Val Ser Val Ser Leu Phe Thr Pro Leu Thr Ala Ala Glu
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Met Ala Pro Tyr Met Lys Arg Leu Ser Arg Gly Gln Thr Val Glu Gly
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Glu Ser Gly Pro Ala Ser Pro Thr Pro Asp Leu Leu Glu Val Leu Ser
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Asp Ile Asp Glu Met Ser Arg Arg Pro Glu Ile Leu Ser Phe Phe
                        535
Ser Thr Asn Leu Gln Arg Leu Met Ser Ser Ala Glu Glu Cys Cys Arg
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Asn Leu Ala Phe Ser Leu Ala Leu Arg Ser Met Gln Asn Ser Pro Ser
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Ile Ala Ala Ala Phe Leu Pro Thr Phe Met Tyr Cys Leu Gly Ser Gln
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Asp Phe Glu Val Val Gln Thr Ala Leu Arg Asn Leu Pro Glu Tyr Ala
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Leu Leu Cys Gln Glu His Ala Ala Val Leu Leu His Arg Ala Phe Leu
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                                            620
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Val Gly Met Tyr Gly Gln Met Asp Pro Ser Ala Gln Ile Ser Glu Ala
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Leu Arg Ile Leu His Met Glu Ala Val Met
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420
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Thr Asn Cys Lys Gln Ala Glu Arg Pro Asn Asn Gln Gln Asn Cys Phe
                            40
Lys Val Cys Asp Trp His Lys Glu Leu Tyr Asp Trp Arg Leu Gly Pro
                                            60
Trp Asn Gln Cys Gln Pro Val Ile Ser Lys Ser Leu Glu Lys Pro Leu
                                        75
                    70
Glu Cys Ile Lys Gly Glu Glu Gly Ile Gln Val Arg Glu Ile Ala Cys
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Ile Gln Lys Asp Lys Asp Ile Pro Ala Glu Asp Ile Ile Cys Glu Tyr
                                105
Phe Glu Pro Lys Pro Leu Leu Glu Gln Ala Cys Leu Ile Pro Cys Gln
                            120
        115
Gln Asp Cys Ile Val Ser Glu Phe Ser Ala Trp Ser Glu Cys Ser Lys
                        135
Thr Cys Gly Ser Gly Leu Gln His Arg Thr Arg His Val Val Ala Pro
                                        155
                    150
Pro Gln Phe Gly Gly Ser Gly Cys Pro Asn Leu Thr Glu Phe Gln Val
                                    170
                165
Cys Gln Ser Ser Pro Cys Glu Ala Glu Glu Leu Arg Tyr Ser Leu His
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Val Gly Pro Trp Ser Thr Cys Ser Met Pro His Ser Arg Gln Val Arq
                            200
                                                 205
Gln Ala Arg Arg Arg Gly Lys Asn Lys Glu Arg Glu Lys Asp Arg Ser
                                            220
                         215
Lys Gly Val Lys Asp Pro Glu Ala Arg Glu Leu Ile Lys Lys Arg
                                         235
                    230
Asn Arg Asn Arg Gln Asn Arg Gln Glu Asn Lys Tyr Trp Asp Ile Gln
                                     250
                 245
 Ile Gly Tyr Gln Thr Arg Glu Val Met Cys Ile Asn Lys Thr Gly Lys
                                 265
 Ala Ala Asp Leu Ser Phe Cys Gln Gln Glu Lys Leu Pro Met Thr Phe
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                                                 285
 Gln Ser Cys Val Ile Thr Lys Glu Cys Gln Val Ser Glu Trp Ser Glu
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295
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Trp Ser Pro Cys Ser Lys Thr Cys His Asp Met Val Ser Pro Ala Gly
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Thr Arg Val Arg Thr Arg Thr Ile Arg Gln Phe Pro Ile Gly Ser Glu
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Lys Glu Cys Pro Glu Phe Glu Glu Lys Glu Pro Cys Leu Ser Gln Gly
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                                345
Asp Gly Val Val Pro Cys Ala Thr Tyr Gly Trp Arg Thr Thr Glu Trp
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Thr Glu Cys Arq Val Asp Pro Leu Leu Ser Gln Gln Asp Lys Arg Arg
                        375
Gly Asn Gln Thr Ala Leu Cys Gly Gly Gly Ile Gln Thr Arg Glu Val
                    390
                                        395
Tyr Cys Val Gln Ala Asn Glu Asn Leu Leu Ser Gln Leu Ser Thr His
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                                    410
Lys Asn Lys Glu Ala Ser Lys Pro Met Asp Leu Lys Leu Cys Thr Gly
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                                425
Pro Ile Pro Asn Thr Thr Gln Leu Cys His Ile Pro Cys Pro Thr Glu
                            440
                                                445
Cvs Glu Val Ser Pro Trp Ser Ala Trp Gly Pro Cys Thr Tyr Glu Asn
                        455
                                            460
Cys Asn Asp Pro Gln Gly Lys Lys Gly Phe Lys Leu Arg Lys Arg Arg
                                        475
                    470
Ile Thr Asn Glu Pro Thr Gly Gly Ser Gly Leu Thr Gly Asn Cys Pro
                                    490
                                                        495
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His Leu Leu Glu Ala Ile Pro Cys Glu Glu Pro Ala Cys Tyr Asp Trp
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Lys Ala Val Arg Leu Gly Asp Cys Glu Pro Asp Asn Gly Lys Glu Cys
                            520
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Gly Pro Gly Thr Gln Val Gln Glu Val Val Cys Ile Asn Ser Asp Gly
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                                            540
Glu Glu Val Asp Arg Gln Leu Cys Arg Asp Ala Ile Phe Pro Ile Pro
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Val Ala Cys Asp Ala Pro Cys Pro Lys Asp Cys Val Leu Ser Thr Trp
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Ser Thr Trp Ser Ser Cys Ser His Thr Cys Ser Gly Lys Thr Thr Glu
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ttataccaat ataaacaatt actcaqqaaa aaaagaaaat aaaaacttgc aagggctaaa
180
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135
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Gly Ala Leu Ser Leu His Leu Pro Glu Gly Arg Asn Ala Val Ser Leu
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                    150
Gln His Arg Arg Asn Thr Ser Glu Lys Lys Ser Ser Arg Lys Val Glu
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Asn Lys Glu Met Glu Tyr Ile Tyr Glu Asn Tyr Tyr Ile
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            180
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gacactecca ggetgagtet entettggtg attetgggeg teatetteat gaatggeaae
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cgtgccagcg aggctgtcct ctgggaggca ctacgcaaga tgggactgcg ccctggggtg
aggcacccat teeteggega tetgaggaag eteateacag atgaetttgt gaagcagaag
tacctggaat acaagaagat ccccaacagc aacccacctg agtatgaatt cctctggggc
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 aa
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<213> Homo sapiens

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Gly Val Ile Phe Met Asn Gly Asn Arg Ala Ser Glu Ala Val Leu Trp
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Glu Ala Leu Arg Lys Met Gly Leu Arg Pro Gly Val Arg His Pro Phe
Leu Gly Asp Leu Arg Lys Leu Ile Thr Asp Asp Phe Val Lys Gln Lys
                    70
Tyr Leu Glu Tyr Lys Lys Ile Pro Asn Ser Asn Pro Pro Glu Tyr Glu
                                    90
Phe Leu Trp Gly Leu Arg Ala Arg His Glu Thr Ser Lys Met Arg Val
                                                     110
                                105
            100
Leu Arg Phe Ile Ala Gln Asn Gln Asn Arg Asp Pro Arg Glu Trp Lys
                            120
Ala His Phe Leu Glu Ala Val Asp Asp Ala Phe Lys Thr Met Asp Val
                        135
Asp Met Ala Glu Glu His Ala Arg Ala Gln Met Arg Ala Gln Met Asn
                                         155
                    150
Ile Gly Asp Glu Ala Leu Ile Gly Arg Trp Ser Trp Asp Asp Ile Gln
                                                         175
                                     170
Val Glu Leu Leu Thr Trp Asp Glu Asp Gly Asp Phe Gly Asp Ala Trp
                                 185
Ala Arg Ile Pro Phe Ala Phe Trp Ala Arg Tyr His Gln Tyr Ile Leu
                                                 205
                             200
        195
Asn Ser Asn Arg Ala Asn Arg Arg Ala Thr Trp Arg Ala Gly Val Ser
                                             220
                         215
Ser Gly Thr Asn Gly Gly Ala Ser Thr Ser Val Leu Asp Gly Pro Ser
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                                         235
Thr Ser Ser Thr Ile Arg Thr Arg Asn Ala Ala Arg Ala Gly Ala Ser
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- <211> 344
- <212> PRT
- <213> Homo sapiens

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                           40
Glu Gln Pro Gly Gln Leu Ile Ser Phe Ser Glu Ala Leu Gln His Phe
                                           60
Gln Thr Val Asp Leu Ser Pro Phe Lys Lys Arg Ile Gln Pro Thr Ile
                   70
                                       75
Arg Arg Thr Gly Leu Ala Ala Leu Arg His Tyr Leu Phe Gly Pro Pro
                                   90
Lys Leu His Gln Arg Leu Arg Glu Glu Arg Asp Leu Val Leu Thr Ile
                               105
Ala Gln Cys Gly Leu Asp Ser Gln Asp Pro Val His Gly Arg Val Leu
                                                125
        115
                            120
Gln Thr Ile Tyr Lys Lys Leu Thr Gly Ser Lys Phe Asp Cys Ala Leu
                       135
His Gly Asn His Trp Glu Asp Leu Gly Phe Gln Gly Ala Asn Pro Ala
                                       155
                    150
Thr Asp Leu Arg Gly Ala Gly Phe Leu Ala Leu Leu His Leu Leu Tyr
                165
                                   170
Leu Val Met Asp Ser Lys Thr Leu Pro Met Ala Gln Glu Ile Phe Arg
                                185
Leu Ser Arg His His Ile Gln Gln Phe Pro Phe Cys Leu Met Ser Val
                            200
Asn Ile Thr His Ile Ala Ile Gln Ala Leu Arg Glu Glu Cys Leu Ser
                                            220
                        215
Arg Glu Cys Asn Arg Gln Gln Lys Val Ile Pro Val Val Asn Ser Phe
                    230
                                        235
Tyr Ala Ala Thr Phe Leu His Leu Ala His Val Trp Arg Thr Gln Arg
                                    250
                245
Lys Thr Ile Ser Asp Ser Gly Phe Val Leu Lys Gly Val Leu Phe Leu
                                265
Leu Gly Arg Pro Arg Leu Asn Ala Gln Cys Pro Arg Ser Arg Glu Pro
                            280
Lys Val Val Ala Arg Leu Val Leu Ala Ala Val Leu Pro His Pro His
                        295
                                            300
Phe Leu Lys Phe Gln Leu Thr Lys Ile Ser Ile Thr His Pro Leu Glu
                    310
                                        315
Ser Ala Ser Ser Pro Phe Ser Ala Leu Thr Val Ala Leu Phe Trp Ser
                325
                                    330
Tyr Thr Tyr Asp Lys His Ile Phe
            340
<210> 4527
<211> 885
<212> DNA
<213> Homo sapiens
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070

<400> 4527

getgeattta ttgtteecag eceggegaga aggtgtteec agaaaggtte ettgggteae ctgcccaccc agccttggct ctgggctgcc atgtccccac gggggcagga gagaggcaca agtcacagte aggcaaggga geetcagegt eetgggeggt ggetgttggg gteeetecag tetteacetg ggaccetegg ccaggetggg acagcateca ggaggegagg etgeatggte cageggtggg tgcaggtggc aacaggtegg egggetgtgc aggttecaaa aggagetete gggttggcac tgggtgagac cagccccggg gccagcaggg gaatgagcgg tggagcaggg 420 ggttgctggg cactggggtg ggccccatct cctgtccttc cctcatggct gctggaaggg 480 cogcetecet ggeteageat cateteagat teegggacte aaacacegte teetegtege tgtccagcga ggccatetec gtggggteet cagtgttgge gaggaggeeg tategeetee gctgaggctt cttcaaccta aacgcccgga tcaggaagta gagcgcggtc aggccgcaga ageccaggat caegtagaag gagegegtea gegeegagee egaegeeeee ggeggaegeg tgtgcgtgct gttgtgtggc gcgcccggct ggctcccgtt cgtcacggcc ggcggcggcg acaacgtgac ctggcggggg cagcggcgag cctcttcggc accgcacggc agcgccgcca gcagcagcgc cagcaggagc agcagcagcg gcggctgcag cacgc 885 <210> 4528 <211> 206 <212> PRT <213> Homo sapiens <400> 4528 Cys Arg Asp Met Ala Ala Phe Ile Val Pro Ser Pro Ala Arg Arg Cys Ser Gln Lys Gly Ser Leu Gly His Leu Pro Thr Gln Pro Trp Leu Trp Ala Ala Met Ser Pro Arg Gly Gln Glu Arg Gly Thr Ser His Ser Gln 50 Ala Arg Glu Pro Gln Arg Pro Gly Arg Trp Leu Leu Gly Ser Leu Gln 70 Ser Ser Pro Gly Thr Leu Gly Gln Ala Gly Thr Ala Ser Arg Arg Arg 85 90 Gly Cys Met Val Gln Arg Trp Val Gln Val Ala Thr Gly Arg Arg Ala 100 105 Val Gln Val Pro Lys Gly Ala Leu Gly Leu Ala Leu Gly Glu Thr Ser 120 Pro Gly Ala Ser Arg Gly Met Ser Gly Gly Ala Gly Gly Cys Trp Ala 140 135 Leu Gly Trp Ala Pro Ser Pro Val Leu Pro Ser Trp Leu Leu Glu Gly

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150
145
Pro Pro Pro Trp Leu Ser Ile Ile Ser Asp Ser Gly Thr Gln Thr Pro
                                    170
               165
Ser Pro Arg Arg Cys Pro Ala Arg Pro Ser Pro Trp Gly Pro Gln Cys
            180
                                185
Trp Arg Gly Gly Arg Ile Ala Ser Ala Glu Ala Ser Ser Thr
                            200
<210> 4529
<211> 546
<212> DNA
<213> Homo sapiens
<400> 4529
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qtqqccgccg cctaagctgc agccgccgga gccgcagaaa caagaggccg agccgtgtcg
aagatggagg agaaaccete agggeeeate eeggacatge tggeeactge agageeeage
tccagtgaga ccgacaagga ggtgttgtcc ccggctgtgc cagctgcagc cccctcctcc
tccatgtcgg aggagccagg ccctgagcag gcagccacac cgccagtggg gaacgtggag
gggctggagg gatgcagcag ggctcctccc cagccccaga cagctgccag tetggccccg
gacccaqccc tggcctgacc agcatagtct ccgggaccag cgaggacctg cggcctccca
qacgacgccc acctccaggg aagcaaatcc cttgctccag ccctggctgc tgcctcagtt
ttcccagcgt ccgtgacctg gcacagcatc tgcgaaccca ctgcccgccg agccctatgc
540
agtctc
546
<210> 4530
<211> 84
<212> PRT
<213> Homo sapiens
<400> 4530
Met Glu Glu Lys Pro Ser Gly Pro Ile Pro Asp Met Leu Ala Thr Ala
Glu Pro Ser Ser Ser Glu Thr Asp Lys Glu Val Leu Ser Pro Ala Val
                                                     30
                                 25
Pro Ala Ala Ala Pro Ser Ser Ser Met Ser Glu Glu Pro Gly Pro Glu
                             40
Gln Ala Ala Thr Pro Pro Val Gly Asn Val Glu Gly Leu Glu Gly Cys
                         55
Ser Arg Ala Pro Pro Gln Pro Gln Thr Ala Ala Ser Leu Ala Pro Asp
                     70
 Pro Ala Leu Ala
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<210> 4531

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<211> 1414
<212> DNA
<213> Homo sapiens
<400> 4531
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geeggteect tgeagggegg tggggeeegg geeetggaee tacteegggg eetgeegegt
gtgagcctgg ccaacttaaa gccgaatccc ggctccaaga aaccggagag aagaccaaga
ggtcggagaa gaggtagaaa atgtggcaga ggccataaag gagaaaggca aagaggaacc
240
eggeceeget tgggetttga gggaggeeag actecatttt acateegaat cecaaaatae
300
gggtttaacg aaggacatag tttcagacgc cagtataagc ctttgagtct caatagactg
360
cagtatetta ttgatttggg tegtgttgat cetagteaac etattgaett aacceagett
gtcaatggga gaggtgtgac catccagcca cttaaaaggg attatggtgt ccagctggtt
gaggagggtg ctgacacctt tacggcaaaa gttaatattg aagtacagtt ggcttcagaa
ctagctattg ctgccattga aaaaaatggt ggtgttgtta ctacagcctt ctatgatcca
agaagtctgg acattgtatg caaacctgtt ccattctttc ttcgtggaca acccattcca
aaaagaatgc ttccaccaga agaactggta ccatattaca ctgatgcaaa gaaccgtggg
tacctggcgg atcctgccaa atttcctgaa gcacgacttg aactcgccag gaagtatggt
780
tatatettae etgatateae taaagatgaa etetteaaaa tgetetgtae taggaaggat
840
ccaaggcaga ttttctttgg tcttgctcca ggatgggtgg tgaatatggc cgataagaaa
atcctaaaac ctacagatga aaatctcctt aagtattata cctcatgaat tcccgtccaa
ggaagcagag ttgttaaaga gtactggaat aggggctgaa ggatctatat tcccttattg
1020
cattttcctt atgtataatt ttccagatgg tgatgttact tttcagtgta ctcatatgtc
tcattttcat ctaaaattaa atggcaggaa acaaggactg catagagaaa ctgagtctgt
gtgggttctg tctcaaagat acaaactccc tgatagtcta tggaaggaaa atgacaacta
ttttagaata tttctagttt gttttttcag tgatcttttc atccaggcct tgttactgtt
acagatcaga atgaaatgca caagtggaat gggattgacc tgtaggcctg ctctgccgag
1320
atgagagcag atggaatgag ttggtgaccc ctcttaatct gtagcctcag ggaaacacgg
ctacccaatg ccaagatggt aaaccctcac gcgt
1414
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<210> 4532
<211> 296
<212> PRT
<213> Homo sapiens
<400> 4532
Met Ala Gly Pro Leu Gln Gly Gly Gly Ala Arg Ala Leu Asp Leu Leu
Arg Gly Leu Pro Arg Val Ser Leu Ala Asn Leu Lys Pro Asn Pro Gly
Ser Lys Lys Pro Glu Arg Arg Pro Arg Gly Arg Arg Gly Arg Lys
                            40
Cys Gly Arg Gly His Lys Gly Glu Arg Gln Arg Gly Thr Arg Pro Arg
                        55
Leu Gly Phe Glu Gly Gly Gln Thr Pro Phe Tyr Ile Arg Ile Pro Lys
                    70
                                        75
Tyr Gly Phe Asn Glu Gly His Ser Phe Arg Arg Gln Tyr Lys Pro Leu
                85
Ser Leu Asn Arg Leu Gln Tyr Leu Ile Asp Leu Gly Arg Val Asp Pro
                                105
Ser Gln Pro Ile Asp Leu Thr Gln Leu Val Asn Gly Arg Gly Val Thr
                            120
Ile Gln Pro Leu Lys Arg Asp Tyr Gly Val Gln Leu Val Glu Glu Gly
                        135
Ala Asp Thr Phe Thr Ala Lys Val Asn Ile Glu Val Gln Leu Ala Ser
                                        155
                    150
Glu Leu Ala Ile Ala Ala Ile Glu Lys Asn Gly Gly Val Val Thr Thr
                165
                                    170
Ala Phe Tyr Asp Pro Arg Ser Leu Asp Ile Val Cys Lys Pro Val Pro
                                 185
Phe Phe Leu Arg Gly Gln Pro Ile Pro Lys Arg Met Leu Pro Pro Glu
                                                 205
                            200
Glu Leu Val Pro Tyr Tyr Thr Asp Ala Lys Asn Arg Gly Tyr Leu Ala
                        215
Asp Pro Ala Lys Phe Pro Glu Ala Arg Leu Glu Leu Ala Arg Lys Tyr
                                         235
                    230
Gly Tyr Ile Leu Pro Asp Ile Thr Lys Asp Glu Leu Phe Lys Met Leu
                                     250
                245
 Cys Thr Arg Lys Asp Pro Arg Gln Ile Phe Phe Gly Leu Ala Pro Gly
                                265
 Trp Val Val Asn Met Ala Asp Lys Lys Ile Leu Lys Pro Thr Asp Glu
                             280
        275
 Asn Leu Leu Lys Tyr Tyr Thr Ser
                         295
    290
 <210> 4533
 <211> 968
 <212> DNA
 <213> Homo sapiens
 <400> 4533
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tttgcacacg tgtgcccctg tccggacgcc ggggctgagg ccgatcgcgt cgggcagcgg
gegeggegge eccgegeage catggactgg etcatgggga agtecaaage caageecaat
ggcaagaagc ccgctgcgga ggagaggaag gcctacctgg agcctgagca caccaaggcc
aggatcaccg acttccagtt caaggagctg gtggtgctgc cccgggagat cgacctcaac
gagtggctgg ccagcaacac aacaacattt ttccaccaca tcaacctgca gtatagcaca
atotoggagt totgcacagg agagacgtgt cagacgatgg cogtgtgcaa cacacagtac
tactggtatg acgagcgggg gaagaaggtc aagtgcacgg ccccacagta cgttgacttc
gtcatgagct ccgtgcagaa gctggtgacg gatgaggacg tgttccccac aaaatacggc
agagaattcc ccageteett tgagteeetg gtgaggaaga tetgeagaca cetgtteeac
qtqctggcac acatctactg ggcccacttc aaggagacgc tggccctgga gctgcacgga
cacttgaaca cgctctacgt ccacttcatc ctctttgctc gggagttcaa cctgctggac
cccaaagaga ccgccatcat ggacgacctc accgaggtgc tatgcagcgg ggccggcggg
gtccacagtg ggggcagtgg ggatggggcc ggcagcgggg gcccgggagc acagaaccac
gtgaaggaga gatgagcccc ccgggccgga caggggcaca cgtgtgcaaa gagacggtgg
tqtqtgttct ctcctgcatc tgcgtgtgca cacatgtgct gggccctctc agacctcacc
960
acacgcgt
968
<210> 4534
<211> 284
<212> PRT
<213> Homo sapiens
<400> 4534
Thr Arg Ala Gln His Met Cys Ala His Ala Asp Ala Gly Glu Asn Thr
His His Arg Leu Phe Ala His Val Cys Pro Cys Pro Asp Ala Gly Ala
Glu Ala Asp Arg Val Gly Gln Arg Ala Arg Arg Pro Arg Ala Ala Met
Asp Trp Leu Met Gly Lys Ser Lys Ala Lys Pro Asn Gly Lys Lys Pro
Ala Ala Glu Glu Arg Lys Ala Tyr Leu Glu Pro Glu His Thr Lys Ala
                                        75
Arg Ile Thr Asp Phe Gln Phe Lys Glu Leu Val Val Leu Pro Arg Glu
                                    90
Ile Asp Leu Asn Glu Trp Leu Ala Ser Asn Thr Thr Thr Phe Phe His
                                 105
His Ile Asn Leu Gln Tyr Ser Thr Ile Ser Glu Phe Cys Thr Gly Glu
```

```
120
        115
Thr Cys Gln Thr Met Ala Val Cys Asn Thr Gln Tyr Tyr Trp Tyr Asp
                        135
Glu Arg Gly Lys Lys Val Lys Cys Thr Ala Pro Gln Tyr Val Asp Phe
                    150
                                        155
Val Met Ser Ser Val Gln Lys Leu Val Thr Asp Glu Asp Val Phe Pro
                165
                                    170
Thr Lys Tyr Gly Arg Glu Phe Pro Ser Ser Phe Glu Ser Leu Val Arg
                                185
                                                     190
Lys Ile Cys Arg His Leu Phe His Val Leu Ala His Ile Tyr Trp Ala
                            200
His Phe Lys Glu Thr Leu Ala Leu Glu Leu His Gly His Leu Asn Thr
                        215
                                            220
Leu Tyr Val His Phe Ile Leu Phe Ala Arg Glu Phe Asn Leu Leu Asp
                                        235
                    230
                                                             240
Pro Lys Glu Thr Ala Ile Met Asp Asp Leu Thr Glu Val Leu Cys Ser
                245
                                    250
Gly Ala Gly Gly Val His Ser Gly Gly Ser Gly Asp Gly Ala Gly Ser
                                265
Gly Gly Pro Gly Ala Gln Asn His Val Lys Glu Arg
                            280
<210> 4535
<211> 473
<212> DNA
<213> Homo sapiens
<400> 4535
cqactttttt ttttttttt tttttgagatg gagtctcgtt ctgtcaccca ggctggagtg
cagtggcatg atcacagete actgcaacet etgeeteeca ggttcaagea gttetetnge
ctcagcctcc cgagtagctg ggattacagg cgtccgccac cacgcccggc taatttttgt
atttttagta gaaacggggt ttcaccatct cggccaggct ggtcttgaac tcctgacctc
atgatecate egecttggee teccaaagtg etgggattae aggeatgage tacegegeee
ggccttggct gcagattaac gggaatacct cccttgggct tcctaggtga cactgtgata
ttcqqtatqa cctcccttqc tctattcctt ggaagaagta caggcactgg tcaagagtgc
cogggaccca cattgcctgg ttttgaatcc cagcacctcc acatgttacg cgt
<210> 4536
<211> 75
<212> PRT
<213> Homo sapiens
<400> 4536
Arg Leu Phe Phe Phe Phe Phe Glu Met Glu Ser Arg Ser Val Thr
                                    10
Gln Ala Gly Val Gln Trp His Asp His Ser Ser Leu Gln Pro Leu Pro
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20
                                25
Pro Arg Phe Lys Gln Phe Ser Xaa Leu Ser Leu Pro Ser Ser Trp Asp
                            40
Tyr Arg Arg Pro Pro Pro Arg Pro Ala Asn Phe Cys Ile Phe Ser Arg
                        55
Asn Gly Val Ser Pro Ser Arg Pro Gly Trp Ser
65
                    70
                                         75
<210> 4537
<211> 2811
<212> DNA
<213> Homo sapiens
<400> 4537
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ctggcttttt acctagcaaa qacaactgaq qctqaqqaaq tctttqtqcc aqttttaaat
ataaaacgtt ctgaactacc tctgcgaggt gacattgtct tctttcttca gaaggttcat
attecagaga gtatettgat ttttegggat gagattgace tecatgcatt ataccagget
ggccaactca ccctcatcct tgtcgaccat catatcttat ccaaaagtga cacaqcccta
gaggagngca gtagcagagg tgctagacca tcgacccatc gagccgaaac actgccctcc
ctgnnccatg tttcagttga gctggtgggg tcctgtgcta ccctggtgac cgagagaatc
ctgcaggggg caccagagat cttggacagg caaactgcag cccttctgca tggaaccatc
atcctggact gtgtcaacat ggaccttaaa attggaaagg caaccccaaa ggacagcaaa
tatgtggaga aactagaggc ccttttccca gacctaccca agagaaatga tatatttgat
tecetacaaa aggeaaagtt tgatgtatea ggactgacea etgageagat getgagaaaa
gaccagaaga ctatctatag acaaggegtc aaggtggcca ttagtgcaat atatatggat
ttggaggeet ttetgeagag gtetaacete ettgeagate teeatgettt etgeeagget
cacagetatg atgtectggt tgccatgact atetttttca acacteacaa tgagecagtg
eggeagttgg ctattttetg tecceatgtg geacteeaaa caacgatetg tgaagteetg
gaacgetece actetecace cetgaagetg acceetgeet caagtaceca cectaacete
960
catgoctate tteaaggeaa caeccaggte tetegaaaga aaettetgee cetgeteeag
1020
gaagccctgt cagcatattt tgactccatg aagatccctt caggacagcc tgagacagca
1080
gatgtgtcca gggagcaagt ggacaaggaa ttggacaggg caagtaactc cctgatttct
1140
ggactgagtc aagatgagga ggaccctccg ctgcccccga cgcccatgaa cagcttggtg
1200
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gatgagtgcc ctctagatca ggggctgcct aaactctctg ctgaggccgt cttcgagaag 1260 tgcagtcaga tctcactgtc acagtctacc acagcctccc tgtccaagaa gtgactgttg agaggegagg aggtagtggg tgaggetace tgaetcaett caaatgeatg ttttgagatg 1380 tttggagatt cagcaattct gtcttcattg ctccaggatc tggtatactg ttctcataaa 1440 actgagagga gaaaaaaagt gaaagaaagc agctgcttta agaatggttt tccacctttt occoctaato totaccaato agacacattt tattatttaa atotgoacot otototattt 1560 tatttqccaq qqqcacqatq tqacatatct gcagtcccag cacagtggga caaaaagaat ttagacccca aaagtgtcct cggcatggat cttgaacaga accagtatct gtcatggaac tgaacattca tcgatggtct ccatgtattc atttattcac ttgttcattc aagtatttat 1740 tgaatacctg cctcaagcta gagagaaaag agagtgcgct ttggaaattt attccagttt tcagcctaca gcagattatc agctcggtga cttttctttc tgccaccatt taggtgatgg tgtttgattc agagatggct gaatttctat tcttagctta ttgtgactgt ttcagatcta gtttgggaac agattagagg ccattgtett etgteetgat caggtggeet ggetgtttet 1980 ttggatccct ctgtcccaga gccacccaga accctgactc ttgagaatca agaaaacacc cagaaaggcc ttaatgacct cataggcact cttccaaaaa gacaacagaa ctggaatgag 2100 aggcctgggt ctgtctcctg ccttagcagg cctatcaatt tcttgtcaat ctctttttt ccttgctcac attaaaagga agcatggagt tctaatgctc ccataaacta tgtattttgg 2220 caagacactt cactactcca ggtctcactt tccccatctg taaaacaggg tttggactag gtgttccctg gtattctgtg atctgcctct tgctgccatt ctttctctcc tctgcttctc tgtatttttc ttctgttatc cctgggggtg ctcaggttca cttgattgtc tgtatttctg tgtggttgta gcaaggactc agcctcatgt agcacgaata ggggtgtggt tcatggcgtg ttgacccagc agagcactcc ctcccactaa cttgttctgc atgtgtagag tctccccatt ttttttaacg caaccettte ceetttttee taccecacag etetgtteca tgtaagttge caacagtttc actgaacagt ggggtatgtg atggttttgg catgacatct tcagtatgag ggggacagtt tgacttcact ttgagggtgt gatgtctgta gctatgtgga aggtaaaaat 2700 agtggtgtga tcatgaacca aaggaattta tgttttgtaa cttgggtact ttattttgca 2760 2811

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<210> 4538
<211> 437
<212> PRT
<213> Homo sapiens
<400> 4538
Xaa Ala Trp His Glu Gly Asn Glu Ala Cys Asp Leu Asp Ser Thr Val
Ser Ala Leu Ala Leu Ala Phe Tyr Leu Ala Lys Thr Thr Glu Ala Glu
                               25
Glu Val Phe Val Pro Val Leu Asn Ile Lys Arg Ser Glu Leu Pro Leu
                           40
                                                45
Arg Gly Asp Ile Val Phe Phe Leu Gln Lys Val His Ile Pro Glu Ser
                        55
Ile Leu Ile Phe Arg Asp Glu Ile Asp Leu His Ala Leu Tyr Gln Ala
                    70
                                        75
Gly Gln Leu Thr Leu Ile Leu Val Asp His His Ile Leu Ser Lys Ser
                85
                                    90
Asp Thr Ala Leu Glu Glu Xaa Ser Ser Arg Gly Ala Arg Pro Ser Thr
                                105
His Arg Ala Glu Thr Leu Pro Ser Leu Xaa His Val Ser Val Glu Leu
                            120
Val Gly Ser Cys Ala Thr Leu Val Thr Glu Arg Ile Leu Gln Gly Ala
                       135
                                           140
Pro Glu Ile Leu Asp Arg Gln Thr Ala Ala Leu Leu His Gly Thr Ile
                    150
                                        155
Ile Leu Asp Cys Val Asn Met Asp Leu Lys Ile Gly Lys Ala Thr Pro
                                    170
                165
Lys Asp Ser Lys Tyr Val Glu Lys Leu Glu Ala Leu Phe Pro Asp Leu
                                185
Pro Lys Arg Asn Asp Ile Phe Asp Ser Leu Gln Lys Ala Lys Phe Asp
                            200
Val Ser Gly Leu Thr Thr Glu Gln Met Leu Arg Lys Asp Gln Lys Thr
                        215
Ile Tyr Arg Gln Gly Val Lys Val Ala Ile Ser Ala Ile Tyr Met Asp
                    230
                                        235
Leu Glu Ala Phe Leu Gln Arg Ser Asn Leu Leu Ala Asp Leu His Ala
                                    250
                245
Phe Cys Gln Ala His Ser Tyr Asp Val Leu Val Ala Met Thr Ile Phe
                               265
Phe Asn Thr His Asn Glu Pro Val Arg Gln Leu Ala Ile Phe Cys Pro
                            280
                                                285
His Val Ala Leu Gln Thr Thr Ile Cys Glu Val Leu Glu Arg Ser His
                        295
                                            300
Ser Pro Pro Leu Lys Leu Thr Pro Ala Ser Ser Thr His Pro Asn Leu
                    310
                                        315
His Ala Tyr Leu Gln Gly Asn Thr Gln Val Ser Arg Lys Lys Leu Leu
                                    330
                325
Pro Leu Leu Gln Glu Ala Leu Ser Ala Tyr Phe Asp Ser Met Lys Ile
                                345
                                                    350
Pro Ser Gly Gln Pro Glu Thr Ala Asp Val Ser Arg Glu Gln Val Asp
                            360
                                                365
Lys Glu Leu Asp Arg Ala Ser Asn Ser Leu Ile Ser Gly Leu Ser Gln
```

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370
                        375
Asp Glu Glu Asp Pro Pro Leu Pro Pro Thr Pro Met Asn Ser Leu Val
                    390
                                        395
Asp Glu Cys Pro Leu Asp Gln Gly Leu Pro Lys Leu Ser Ala Glu Ala
                                    410
                405
Val Phe Glu Lys Cys Ser Gln Ile Ser Leu Ser Gln Ser Thr Thr Ala
            420
                                425
Ser Leu Ser Lys Lys
        435
<210> 4539
<211> 331
<212> DNA
<213> Homo sapiens
<400> 4539
gtgcacggag gaaagtctca tgagcagcct gaatgggggc tctgttcctt ctgagctgga
tgggctggac tccgagaaag accagaagcc tgggggaaaa ccaaagggta atcaatgaac
tcacctggaa actccagcaa gagcagaggc aggtggagga gctgaggatg cagcttcaga
agcagaaaag gaataactgt tcagagaaga agccgctgcc tttcctggct gcctccatca
aqcaaqaaqa qqctgtctcc agctgtcctt ttgcatccca agtacctgtg aaaagacaaa
geageagete aaagtgteae eeaceggett g
331
<210> 4540
<211> 99
<212> PRT
<213> Homo sapiens
<400> 4540
Met Gly Ala Leu Phe Leu Leu Ser Trp Met Gly Trp Thr Pro Arg Lys
                                    10
Thr Arg Ser Leu Gly Glu Asn Gln Arg Val Ile Asn Glu Leu Thr Trp
Lys Leu Gln Gln Glu Gln Arg Gln Val Glu Glu Leu Arg Met Gln Leu
                            40
Gln Lys Gln Lys Arg Asn Asn Cys Ser Glu Lys Lys Pro Leu Pro Phe
Leu Ala Ala Ser Ile Lys Gln Glu Glu Ala Val Ser Ser Cys Pro Phe
                    70
                                         75
Ala Ser Gln Val Pro Val Lys Arg Gln Ser Ser Ser Lys Cys His
                                                         95
                                     90
Pro Pro Ala
<210> 4541
<211> 452
<212> DNA
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<213> Homo sapiens

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<400> 4541
actagicace tettetatea gaigateate iggaleatat tettitagat taataatgge
cacaggcaga tccagggatg taactgcttc agcaagaact gttgcgaatc ccttcgctgt
tccagtctga gaaccataaa aaatcttcac tccagacaca aagatgtctt tctcttgaag
ggagacataa ccatttgtca tcaaatcctg agctgctttt ggaacagatt tttcctgtaa
gttettgece tgcgtettga tgacaatetg gacacaaate caaaggetaa tgetaacage
aaaqcccaaa taaatgtaaa acctgtttat ccacaatgat attaaaggtg agaagaggtc
ccatqtatcc qcagagggat ccatcctcct cagagccgac aggagactag gatctcggac
ctggagagcc cgatgattcg cactggtact gc
452
<210> 4542
<211> 128
<212> PRT
<213> Homo sapiens
<400> 4542
Met Asp Pro Ser Ala Asp Thr Trp Asp Leu Phe Ser Pro Leu Ile Ser
Leu Trp Ile Asn Arg Phe Tyr Ile Tyr Leu Gly Phe Ala Val Ser Ile
                                25
Ser Leu Trp Ile Cys Val Gln Ile Val Ile Lys Thr Gln Gly Lys Asn
        35
Leu Gln Glu Lys Ser Val Pro Lys Ala Ala Gln Asp Leu Met Thr Asn
                        55
Gly Tyr Val Ser Leu Gln Glu Lys Asp Ile Phe Val Ser Gly Val Lys
                    70
                                        75
Ile Phe Tyr Gly Ser Gln Thr Gly Thr Ala Lys Gly Phe Ala Thr Val
                                    90
Leu Ala Glu Ala Val Thr Ser Leu Asp Leu Pro Val Ala Ile Ile Asn
                                105
Leu Lys Glu Tyr Asp Pro Asp Asp His Leu Ile Glu Glu Val Thr Ser
                            120
                                                 125
        115
<210> 4543
<211> 815
<212> DNA
<213> Homo sapiens
<400> 4543
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agggaggagg gagagcgagt cactgcaggt ccctggcctg cggctccgcc gtggctgcct
gaggecege geaccaatge tttgcaettt geetegeeeg acaecetgeg ggecagaget
180
```

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Thr	Val	Ile		Glu	Arg	Leu	His		Arg	GIY	Ala	Met	430	Pne	мта
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Thr	Glu		Ala	Leu	Ala	ALA	440	Ala	GIII	Giu	1111	445	Arg		Deu
		435	Th	7 ~~	Glu	Ma I		Glu	Δen	TIA	Tvr		Asp	Trp	Gln
cys	450	Ala	IYL	ALG	GIU	455	AIG	Oru	лор		460				
C1 n		uie	Gln	Glu	Ala		Len	Leu	Leu	Gln		Arq	Ala	Gln	Ala
465	ALG		01		470					475		-			480
T.011	Gln	Gln	Va 1	Tvr	Asn	Glu	Met	Glu	Gln	Asp	Leu	Arg	Leu	Leu	Gly
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Ala	Thr	Ala	Ile	Glu	Asp	Arg	Leu	Gln	Asp	Gly	Val	Pro	Glu	Thr	Ile
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Lys	Cys	Leu	Lys	Lys	Ser	Asn	Ile	Lys	Ile	Trp	Val	Leu	Thr	Gly	Asp
		515					520					525		_	_
Lys	Gln	Glu	Thr	Ala	Val		Ile	Gly	Phe	Ala	Cys	Glu	Leu	Leu	Ser
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	Asn	Met	Leu	Ile	Leu	Glu	GIu	Lys	GIU	555	ser	Arg	ire	Leu	560
545	_	_			550 Ser			T 011	T 011		A ror	Glu	Sar	T.011	
Thr	Tyr	Trp	Glu			Asn	Asn	Leu	570	1111	Arg	GIU	261	575	261
~~				565	Leu	17-1	T10	Aen		Aen	Dhe	Len	Asp		Leu
GIn	vaı	Lys	580		Leu	vai	IIe	585	OLY	АБР			590	-1-	
T 411	t/a1	Ser			Lys	Glu	Pro		Ala	Leu	Ala	Gln	Asn	Val	Asn
Dea	*41	595		5	-3-		600					605			
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625					630					635					640
Ala	Ala	Pro	Pro		Gln	Asp	Ser	Arg			Arg	Ser	Ser	Glu	Val
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Leu	Gln	Glu			Phe	Val	Asp			Ser	Lys	Cys	GIN	ATA	vai
			660	٠		_		665				T1.	670		T 011
Ile	Cys			Val	Thr	Pro			Lys	Ата	Leu	685	val	Ala	Leu
	_	675			Gln	**- 1	680			71.	т1.			Glv	7.1 a
Val			Tyr	HIS	GIN	695		Int	Leu	мта	700	GIY	мэр	Gry	ALU
	690	71.	. 7.00	Mot	Ile			712	Aen	Va1			Glv	Leu	Ala
705		i iie	ASI	nec	710		1111	ALG	ADP	715					720
011	Gla	Glu	G1v	, Met	Gln		Va1	Gln	Asn			Phe	Val	Leu	Gly
GLY	GII	. 010	. 017	725					730		•			735	-
Gl n	Phe	Cvs	Phe		Glr	Ara	r Leu	Leu	Leu	Val	His	Gly	Arg	Trp	Ser
			740	)				745					750		
Tyr	Val	Arg	, Ile	Cys	Lys	Phe	Leu	Arg	Tyr	Phe	Phe	Tyr	Lys	Ser	Met
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Leu Trp Ile Ser Arg Asp Thr Ala Gly Pro Ala Ser Phe Ser Asp His
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Gln Ser Phe Ala Val Val Ala Leu Ser Cys Leu Leu Ser Ile Thr
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Cys Ser Gln Gln Gly Arg Gln Gly Arg Ala Pro Arg Arg Asp Pro Thr
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Gln Arg Thr Trp Glu Ser Gly Cys Gln Arg Trp Ala Ala Gly Arg Ala
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Glu Ile Arg Pro Trp Phe Thr Pro Arg Ser Ile Tyr Met Glu Ala Ser
Thr Val Asp Cys Asn Asp Leu Gly Leu Leu Thr Phe Pro Ala Arg Leu
Pro Ala Asn Thr Gln Ile Leu Leu Leu Gln Thr Asn Asn Ile Ala Lys
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Ile Glu Tyr Ser Thr Asp Phe Pro Val Asn Leu Thr Gly Leu Asp Leu
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Ser Gln Asn Asn Leu Ser Ser Val Thr Asn Ile Asn Val Lys Lys Met
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Pro Gln Leu Leu Ser Val Tyr Leu Glu Glu Asn Lys Leu Thr Glu Leu
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                            120
Pro Glu Lys Cys Leu Ser Glu Leu Ser Asn Leu Gln Glu Leu Tyr Ile
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                                            140
Asn His Asn Leu Leu Ser Thr Ile Ser Pro Gly Ala Phe Ile Gly Leu
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His Asn Leu Leu Arg Leu His Leu Asn Ser Asn Arg Leu Gln Met Ile
                                    170
Asn Ser Lys Trp Phe Asp Ala Leu Pro Asn Leu Glu Ile Leu Met Ile
                                185
            180
Gly Glu Asn Pro Ile Ile Arg Ile Lys Asp Met Asn Phe Lys Pro Leu
                                                205
                            200
Ile Asn Leu Arg Ser Leu Val Ile Ala Gly Ile Asn Leu Thr Glu Ile
                        215
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Pro Asp Asn Ala Leu Val Gly Leu Glu Asn Leu Glu Ser Ile Ser Phe
                                        235
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Tyr Asp Asn Arg Leu Ile Lys Val Pro His Val Ala Leu Gln Lys Val
                                    250
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Val Asn Leu Lys Phe Leu Asp Leu Asn Lys Asn Pro Ile Asn Arg Ile
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Arg Arg Gly Asp Phe Ser Asn Met Leu His Leu Lys Glu Leu Gly Ile
                                                 285
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Asn Asn Met Pro Glu Leu Ile Ser Ile Asp Ser Leu Ala Val Asp Asn
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 Leu Pro Asp Leu Arg Lys Ile Glu Ala Thr Asn Asn Pro Arg Leu Ser
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 Tyr Ile His Pro Asn Ala Phe Phe Arg Leu Pro Lys Leu Glu Ser Leu
                                     330
Met Leu Asn Ser Asn Ala Leu Ser Ala Leu Tyr His Gly Thr Ile Glu
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Cvs Asp Cvs Val Ile Arg Trp Met Asn Met Asn Lys Thr Asn Ile Arg
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Phe Met Glu Pro Asp Ser Leu Phe Cys Val Asp Pro Pro Glu Phe Gln
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Gly Gln Asn Val Arg Gln Val His Phe Arg Asp Met Met Glu Ile Cys
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Leu Pro Leu Ile Ala Pro Glu Ser Phe Pro Ser Asn Leu Asn Val Glu
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Ala Gly Ser Tyr Val Ser Phe His Cys Arg Ala Thr Ala Glu Pro Gln
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Asn Gly Val Thr Pro Lys Glu Gly Gly Leu Tyr Thr Cys Ile Ala Thr
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Asn Leu Val Gly Ala Asp Leu Lys Ser Val Met Ile Lys Val Asp Gly
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Ser Phe Pro Gln Asp Asn Asn Gly Ser Leu Asn Ile Lys Ile Arg Asp
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Ile Gln Ala Asn Ser Val Leu Val Ser Trp Lys Ala Ser Ser Lys Ile
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Leu Lys Ser Ser Val Lys Trp Thr Ala Phe Val Lys Thr Glu Asn Ser
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His Ala Ala Gln Ser Ala Arg Ile Pro Ser Asp Val Lys Val Tyr Asn
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Leu Thr His Leu Asn Pro Ser Thr Glu Tyr Lys Ile Cys Ile Asp Ile
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Pro Thr Ile Tyr Gln Lys Asn Arg Lys Lys Cys Val Asn Val Thr Thr
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Lys Gly Leu His Pro Asp Gln Lys Glu Tyr Glu Lys Asn Asn Thr Thr
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Thr Leu Met Ala Cys Leu Gly Gly Leu Leu Gly Ile Ile Gly Val Ile
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Cys Leu Ile Ser Cys Leu Ser Pro Glu Met Asn Cys Asp Gly Gly His
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Ser Tyr Val Arg Asn Tyr Leu Gln Lys Pro Thr Phe Ala Leu Gly Glu
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Leu Tyr Pro Pro Leu Ile Asn Leu Trp Glu Ala Gly Lys Glu Lys Ser
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Glu Thr Ser Arg Ala Phe Leu Pro Pro Pro Ser Asp Val Arg Val Arg
Ser Cys Leu Tyr His Trp Ser Ala Thr Ala His Leu Pro Pro Leu Ser
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Lys Lys Pro Pro Cys Thr Ile Ser His Leu Arg Pro Leu Leu Gly Leu
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Pro Pro Pro Ser Asp Leu His Ile Pro Ser Ala Ala Thr Leu Gly Pro
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Cys Met His Trp Pro Pro Pro Ser Asp Ala Pro Cys Thr Ile Ser Leu
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Ser Phe Arg Lys Ile Gly Asn Arg Arg Arg Gln Glu Arg Phe Trp Tyr
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90
Cys Arg Leu Ala Leu Asn His Lys Val Leu His Tyr Gly Asp Leu Asp
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Asp Asn Pro Gln Gly Glu Val Thr Phe Glu Ser Leu Gln Glu Lys Ile
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                             120
                                                 125
Pro Val Ala Asp Ile Lys Ala Ile Val Thr Gly Lys Asp Cys Pro His
    130
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                                             140
Met Lys Glu Lys Ser Ala Leu Lys Gln Asn Lys Glu Val Leu Glu Leu
                    150
                                         155
Ala Phe Ser Ile Leu Tyr Asp Pro Asp Glu Thr Leu Asn Phe Ile Ala
                                     170
                165
Pro Asn Lys Tyr Glu Tyr Cys Ile Trp Ile Asp Gly Leu Ser Ala Leu
            180
                                185
                                                     190
Leu Gly Lys Asp Met Ser Ser Glu Leu Thr Lys Ser Asp Leu Asp Thr
        195
                             200
                                                 205
Leu Leu Ser Met Glu Met Lys Leu Arg Leu Leu Asp Leu Glu Asn Ile
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                        215
                                             220
Gln Ile Pro Glu Ala Pro Pro Pro Ile Pro Lys Glu Pro Ser Ser Tyr
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225
                                                              240
Asp Phe Val Tyr His Tyr Gly
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cacattgtgg tectgaggea cacaggaat ggggeoacet getgacaca ttgtgacgga 300
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accttaggaag agagacaat gattagcatt tatggcattg ggtgcacca acttegtata 720
ggacegtact ectggacac atttecacat gtggattte ggttgcacca agagaacaagaaaag

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caaatactaq aqaatettte caetteqeet etqqetqaqe caececaett tgttgaacat
attagateta cettgatgtt tttaaaaaaaa cacccatete cageteacae actgttttet
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Gly Asp Leu Val Arg Ala His Pro Pro Leu Glu Glu Arg Ala Arg Leu
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Val Gln Gln Arg Glu Leu Ala Val Thr Ser Pro Lys Asp Gly Ser Ile
                       55
Ser Ile Leu Gly Ser Asp Asp Ala Thr Thr Cys His Ile Val Val Leu
                                      75
Arg His Thr Gly Asn Gly Ala Thr Cys Leu Thr His Cys Asp Gly Thr
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Asp Thr Lys Ala Glu Val Pro Leu Ile Met Asn Ser Ile Lys Ser Phe
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Ser Asp His Ala Gln Cvs Gly Arg
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Leu Leu Gly Pro Gly Glu Thr Val Leu Arg Gln Lys Leu Gly Val Gln
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Gly Gly Pro Arg Val Arg His Cys Gly Glu Gly Asn Ala Gly Glu Ser
                   70
Gly Pro Thr Leu Gln Leu Gly Thr Arg Gly Arg Lys Gln Arg Gly Gln
Ala Ser Val Pro Leu Pro Gln Glu Gln Thr Ser Gly Pro Gln Glu Gly
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Leu Gln Ala Ala Arg Ser Leu Pro Ser Ala Gly Gly Ser Arg Gly Arg
Lys Gly Trp Arg Ala Ala Gly Arg Gln Pro Ser Thr Arg
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Lys Thr Gln Gln Asn Arg Lys Leu Thr Asp Phe Tyr Pro Val Arg Arg
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Ser Ser Arg Lys Ser Lys Ala Glu Leu Gln Ser Glu Glu Arg Lys Arg
Ile Asp Glu Leu Ile Glu Ser Gly Lys Glu Glu Gly Met Lys Ile Asp
                        55
    50
Leu Ile Asp Gly Lys Gly Arg Gly Val Ile Ala Thr Lys Gln Phe Ser
                                                             Ωn
                    70
65
Arg Gly Asp Phe Val Val Glu Tyr His Gly Asp Leu Ile Glu Ile Thr
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Asp Ala Lys Lys Arg Glu Ala Leu Tyr Ala Gln Asp Pro Ser Thr Gly
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Cys Tyr Met Tyr Tyr Phe Gln Tyr Leu Ser Lys Thr Tyr Trp
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Gly Pro Thr Gly Pro Lys Gly Asp Ala Gly Ser Arg Gly Pro Met Gly
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Met Arg Gly Pro Pro Gly Pro Gln Gly Pro Pro Gly Ser Pro Gly Arg
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Ala Gly Ala Val Gly Thr Pro Gly Lys Arg Gly Pro Ser Gly Pro Gln
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Gly Leu Leu Gly Pro Pro Gly Pro Pro Ala Pro Val Gly Pro Pro His
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                    70
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Glu Thr Asn Pro Phe Thr Arg
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720
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960
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Gln Ala Ala Leu His Leu Leu Gln Pro Leu Gly His Val Ala Arq Glu
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Pro Ala Arg His Val Ala Thr Ala Gln Gly Glu Val Leu Pro Pro Gly
Gly Leu Gly Gly Ala Ala Gln Arg Ala Arg Gly Gln Ser His Gly Gly
                        55
Thr Val Pro Gly Asn Ala Pro Ala Ala Asp Leu Leu Ala Leu Ser Pro
Arg Leu Glu Arg Ser Gly Thr Ile Ser Thr His Cys Lys Leu Arg Leu
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Pro Gly Ser Arg His Ser Pro Ala Ser Ala Ser
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420
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<211> 1007

<213> Homo sapiens

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Thr	Ile		Asp	Ala	Gln			Asp	Cys	Tyr			Pro	Met	Lys
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Ser	Gln			Arg	Lys	Glu	Val 520		Ala	Gly	Pro	Gly 525		Gln	Gln
Gly	Asp	515 Ser	Tyr	Leu	Arg			Ser	Asp	Ser			Asp	Gln	Ser
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545	Val	Dro	cor	Car		T.011	Dro	Gln	Thr		Glu	Gln	Glu	Laze	
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Leu	Arg	His	His		Glu	Thr	Leu	Thr		Ser	Pro	Cys	Arq		
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Leu	Ser	Ile	Ser	Thr	Gln	Phe	Leu	Ser	Ser	Leu	Gln	Lys	Ala	Ser	Arg
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Phe 625	Thr	His	Thr	Phe	Pro 630	Pro	Arg	Ala	Thr	Gln 635	Сув	Leu	Val	Lys	Ser 640
Pro	Glu	Val	Lys	Leu 645	Met	Asp	Arg	Gly	Gly 650	Ser	Gln	Pro	Arg	Ala 655	Gly
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Met	Glu	Ala	Thr	Ala 725	Ser	Ser	Arg	Ala	Arg 730	Ile	Ser	Arg	Ser	Ile 735	Ser
Leu	Gly	Asp	Ser	Glu	Gly	Pro	Ile	Val 745	Ala	Thr	Leu	Ala	Gln 750	Pro	Leu
Arg	Arg	Pro 755	Ser	Ser	Val	Gly	Glu 760	Leu	Ala	Ser	Leu	Gly 765	Gln	Glu	Leu
Gln	Ala 770		Thr	Thr	Ala	Thr		Pro	Ser	Leu	Asp 780		Glu	Gly	Gln
al.	Pro	nla	T.011	Arm	Car		Glv	Aen	uic	Glu		Arm	ala	Aen	T.011
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Asp	Thr	Gln	Pro	Gly	Val	Thr	Val	Pro 825	Ala	Val	Ser	Phe	Pro 830	Ala	Pro
Ser	Pro	Val		Glu	Ser	Ala	Leu		Leu	His	Gly	Ser		Phe	Arg

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Leu Glu Ile Asp His Arg Gln Gln Gln His Thr Asn Asp Lys Lys
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Leu Ala Lys Lys Glu Ala Lys Glu Arg Lys Lys Arg Glu Lys Met Gly
Trp Gly Glu Glu Tyr Met Gly Tyr Thr Asn Thr Asp Asn Pro Phe Gly
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Asp Asn Asn Leu Leu Gly Thr Phe Ile Trp Asn Lys Ala Leu Glu Lys
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Lys Gly Ile Ser His Leu Glu Glu Lys Glu Leu Lys Glu Arg Asn Lys
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Gln Glu Asp Asn Phe His Leu Gln Gln Ala Lys Leu Arg Ser Lys Ile
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Ile Ser Ala Glu Asp Asp Asp Leu Ala Gly Glu Met His Glu Pro Tyr
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                                                205
Thr Phe Leu Asn Gly Leu Thr Val Ala Asp Met Glu Asp Leu Leu Glu
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Asp Ile Gln Val Tyr Met Glu Leu Glu Gln Gly Lys Asn Ala Asp Phe
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Trp Arg Asp Met Thr Thr Ile Thr Glu Asp Glu Ile Ser Lys Leu Arg
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Lys Leu Glu Ala Ser Gly Lys Gly Pro Gly Glu Arg Arg Glu Gly Val
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Asn Ala Ser Val Ser Ser Asp Val Gln Ser Val Phe Lys Gly Lys Thr
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Tyr Asn Gln Leu Gln Val Ile Phe Gln Gly Ile Glu Gly Lys Ile Arg
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Ala Gly Gly Pro Asn Leu Asp Met Gly Tyr Trp Glu Ser Leu Leu Gln
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Gln Leu Arg Ala His Met Ala Arg Ala Arg Leu Arg Glu Arg His Gln
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Gln Leu Leu Gln Ile Arg Gln Asp Val Glu Ser Cys Tyr Phe Ala Ala
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Ala Asp Leu Ala Leu Gln Met Pro Ser Trp Lys Gly Cys Val Gln Thr
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Glu Ile Leu Thr Val Leu Pro Glu Glu Val His Ser Arg Ser Leu Arg
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Ile Gly Ala Asn Arg Arg Thr Glu Ile Ile Glu Asp Leu Ala Phe Tyr
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Ser Ser Thr Val Val Ser Leu Leu Met Thr Cys Val Glu Lys Ala Gly
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Thr Asp Glu Lys Met Leu Met Lys Val Phe Arg Cys Leu Gly Ser Trp
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                                                205
Phe Asn Leu Gly Val Leu Asp Ser Asn Phe Met Ala Asn Asn Lys Leu
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Leu Ala Leu Leu Phe Glu Val Leu Gln Gln Asp Lvs Thr Ser Ser Asn
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Leu	Arg	Thr	Leu	Glu 325	Leu	Leu	Leu	Ile	Cys 330	Ala	Gly	His	Pro	Gln 335	Tyr
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			500					505					510		
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	530					535					540			Glu	
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Thr	610		Leu	Asp	ALC	615	, WIG	. vai	. 116	· FIIC	620				
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025 T14	Trr	Pro	val	Leu	Set	Glu	ı Thı	Leu	Asr			Arg	Ala	Asp	Asn
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Leu Leu Asp Met Leu Gln Ala Leu Cys Ile Pro Thr Phe Gln Leu Leu
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Phe Arg Leu Ala Thr Arg Phe Ile Gln Arg Ser Pro Val Thr Leu Leu
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                                             780
Thr Leu Asp His Arg Asp Ala Asn Cys Ser Val Met Arg Phe Leu Arg
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Asp Leu Ile His Thr Gly Val Ala Asn Asp His Glu Glu Asp Phe Glu
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Arg Pro Thr Phe Cys Arg Trp Leu Glu Asn Ser Leu Lys Gly Leu Pro
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                                         875
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<212> DNA

<213> Homo sapiens

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Arg Ile Pro Ser Leu Gln Gln Leu His Leu Gln Arg Asn Ala Leu Cys
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                                        75
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His Gln His Leu Lys Thr Leu Leu Leu Glu Arg Asn Pro Ile Lys Met
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Leu Pro Arg Asn Pro Thr Ser Gln Glu Ala Pro Pro Val Arg Glu Met
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Thr Leu Arg Asp Leu Pro Ser Pro Gly Leu Glu Leu Ser Gly Asp His
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Ala Ser Asn Gln Gly Ala Val Asn Ala Gln Asp Pro Glu Gly Ala Val
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Met Lys Glu Lys Ala Ser Phe Leu Pro Pro Val Glu Lys Pro Asp Leu
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Ser Glu Leu Arg Lys Ser Ala Asp Ser Ser Glu Asn Trp Pro Ser Glu
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Glu Glu Ile Arg Arg Phe Trp Lys Leu Arg Gln Glu Ile Val Glu His
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Val Lys Ala Asp Val Leu Gly Asp Gln Leu Leu Thr Arg Glu Leu Pro
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Pro Asn Leu Lys Ala Ala Leu Asn Ile Glu Lys Glu Leu Pro Lys Pro
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Arg His Val Phe Arg Arg Lys Thr Ala Ser Ser Arg Ser Ile Leu Pro
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Asp Leu Leu Ser Pro Tyr Gln Met Ala Ile Arg Ala Lys Arg Leu Glu
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Glu Ser Arg Ala Ala Ala Leu Arg Glu Leu Gln Glu Lys Gln Ala Leu
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Met Glu Gln Gln Arg Arg Glu Lys Arg Ala Leu Gln Glu Trp Arg Glu
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Arg Ala Gln Arg Met Arg Lys Arg Lys Glu Glu Leu Ser Lys Leu Leu
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Pro Pro Arg Arg Ser Met Val Ala Ser Lys Ile Pro Ser Ala Thr Asp
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Gln Glu Arg Asn Leu Glu Glu Lys Ile Lys Gln His Val Leu Gln Met
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Lys Ala Ala Glu Asp Leu Glu Ile Ala Thr Glu Leu Gln Asp Glu Val
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Gln Pro Leu Asp Lys Lys Ala Ala Val Ser Trp Leu Thr Pro Ala Pro
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Ser Lys Lys Ala Asp Ser Val Ala Ala Lys Val Asp Leu Leu Gly Glu
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Phe Gln Ser Ala Leu Pro Lys Ile Asn Ser His Pro Thr Arg Ser Gln
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Lys Lys Ser Ser Gln Lys Lys Ser Ser Lys Lys Asn His Pro Gln Lys
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Asn Ala Pro Gln Asn Ser Thr Gln Ala His Ser Glu Asn Lys Cys Ser
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Pro Pro Cys His Ile Val Asp Tyr Arg Thr Arg Trp Ser Gly Ile Arg
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Ile Leu Lys Ile Leu Thr Gly Lys Ile Val Val Gly His Ala Ile His
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Asn Asp Phe Lys Ala Leu Gln Tyr Phe His Pro Lys Ser Leu Thr Arq
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Asp Thr Ser His Ile Pro Pro Leu Asn Arg Lys Ala Asp Cys Pro Glu
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Asn Ala Thr Met Ser Leu Lys His Leu Thr Lys Lys Leu Leu Asn Arg
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Asp Ile Gln Val Gly Lys Ser Gly His Ser Ser Val Glu Asp Ala Gln
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Gly Phe Arg Pro Gly Met Arg Cys Gly Gly Ser Ser Leu Gly Arg Thr
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Cys Cys Ser Pro Thr Arg Arg Ala Cys Val Val Ser Arg Ala Val Thr
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Thr Val Thr Ser Lys Val Ala Pro Ser Trp Pro Glu Ser His Ser Ser
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Ala Asp Ser Ala Ser Leu Ala Lys Lys Lys Pro Leu Phe Ile Thr Thr
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Mal.															
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Arg Lys 465 Gln	Phe Ser 450 Cys	Gly 435 Glu Ala Ile	420 Val Thr Lys Pro	Cys Glu Gly Gly 485	Leu Glu Gln 470	Asp Met 455 Ser Ala	Cys 440 Gly His Leu	425 Tyr Asp Glu Tyr Lys	Arg Glu Pro Asn 490	Leu Glu Glu 475 Ile	Arg Val 460 Asn Gly	Lys 445 Phe Leu Asp	430 Ser Ser Met Met	Arg Trp Pro Val 495	Pro Leu Thr 480 His
Arg Lys 465 Gln Ala	Phe Ser 450 Cys Ile Ala	Gly 435 Glu Ala Ile Arg	420 Val Thr Lys Pro Gly 500	Cys Glu Gly Gly 485 Lys	Leu Glu Gln 470 Thr	Asp Met 455 Ser Ala Gly	Cys 440 Gly His Leu Ile	425 Tyr Asp Glu Tyr Lys 505	Arg Glu Pro Asn 490 Ala	Leu Glu Glu 475 Ile Asn	Arg Val 460 Asn Gly Cys	Lys 445 Phe Leu Asp	430 Ser Ser Met Met Cys 510	Arg Trp Pro Val 495 Ile	Pro Leu Thr 480 His Ser
Arg Lys 465 Gln Ala	Phe Ser 450 Cys Ile Ala	Gly 435 Glu Ala Ile Arg	420 Val Thr Lys Pro Gly 500	Cys Glu Gly Gly 485 Lys	Leu Glu Gln 470 Thr	Asp Met 455 Ser Ala Gly	Cys 440 Gly His Leu Ile Arg	425 Tyr Asp Glu Tyr Lys 505	Arg Glu Pro Asn 490 Ala	Leu Glu Glu 475 Ile Asn	Arg Val 460 Asn Gly Cys	Lys 445 Phe Leu Asp Pro	430 Ser Ser Met Met Cys 510	Arg Trp Pro Val 495 Ile	Pro Leu Thr 480 His Ser
Arg Lys 465 Gln Ala Arg	Phe Ser 450 Cys Ile Ala Gln	Gly 435 Glu Ala Ile Arg Asn 515	420 Val Thr Lys Pro Gly 500 Lys	Glu Gly Gly 485 Lys Ser	Glu Gln 470 Thr Trp Val	Asp Met 455 Ser Ala Gly Leu	Cys 440 Gly His Leu Ile Arg 520	ASP Glu Tyr Lys 505 Pro	Arg Glu Pro Asn 490 Ala Ala	Glu Glu 475 Ile Asn Val	Arg Val 460 Asn Gly Cys Thr	Lys 445 Phe Leu Asp Pro Asn 525	A30 Ser Ser Met Met Cys 510 Gly	Arg Trp Pro Val 495 Ile Met	Pro Leu Thr 480 His Ser
Arg Lys 465 Gln Ala Arg	Phe Ser 450 Cys Ile Ala Gln Leu	Gly 435 Glu Ala Ile Arg Asn 515	420 Val Thr Lys Pro Gly 500 Lys	Glu Gly Gly 485 Lys Ser	Leu Glu Gln 470 Thr	Asp Met 455 Ser Ala Gly Leu Pro	Cys 440 Gly His Leu Ile Arg 520	ASP Glu Tyr Lys 505 Pro	Arg Glu Pro Asn 490 Ala Ala	Glu Glu 475 Ile Asn Val	Arg Val 460 Asn Gly Cys Thr	Lys 445 Phe Leu Asp Pro Asn 525	A30 Ser Ser Met Met Cys 510 Gly	Arg Trp Pro Val 495 Ile Met	Pro Leu Thr 480 His Ser
Arg Lys 465 Gln Ala Arg	Phe Ser 450 Cys Ile Ala Gln Leu 530	Gly 435 Glu Ala Ile Arg Asn 515 Pro	420 Val Thr Lys Pro Gly 500 Lys Ser	Cys Glu Gly 485 Lys Ser Ile	Leu Glu Gln 470 Thr Trp Val	Asp Met 455 Ser Ala Gly Leu Pro 535	Cys 440 Gly His Leu Ile Arg 520 Ser	Asp Glu Tyr Lys 505 Pro	Arg Glu Pro Asn 490 Ala Ala Ser	Glu Glu 475 Ile Asn Val Ser	Arg Val 460 Asn Gly Cys Thr Gly 540	Lys 445 Phe Leu Asp Pro Asn 525 Asn	A30 Ser Ser Met Cys 510 Gly	Arg Trp Pro Val 495 Ile Met Thr	Pro Leu Thr 480 His Ser Ser
Arg Lys 465 Gln Ala Arg Gln	Phe Ser 450 Cys Ile Ala Gln Leu 530	Gly 435 Glu Ala Ile Arg Asn 515 Pro	420 Val Thr Lys Pro Gly 500 Lys Ser	Cys Glu Gly 485 Lys Ser Ile	Glu Gln 470 Thr Trp Val	Asp Met 455 Ser Ala Gly Leu Pro 535 Pro	Cys 440 Gly His Leu Ile Arg 520 Ser	Asp Glu Tyr Lys 505 Pro	Arg Glu Pro Asn 490 Ala Ala Ser	Glu Glu 475 Ile Asn Val Ser	Arg Val 460 Asn Gly Cys Thr Gly 540	Lys 445 Phe Leu Asp Pro Asn 525 Asn	A30 Ser Ser Met Cys 510 Gly	Arg Trp Pro Val 495 Ile Met Thr	Pro Leu Thr 480 His Ser Ser
Arg Lys 465 Gln Ala Arg Gln Phe 545	Phe Ser 450 Cys Ile Ala Gln Leu 530 Ser	Gly 435 Glu Ala Ile Arg Asn 515 Pro	420 Val Thr Lys Pro Gly 500 Lys Ser	Glu Gly Gly 485 Lys Ser Ile Gly	Leu Glu Gln 470 Thr Trp Val Asn Gly 550	Asp Met 455 Ser Ala Gly Leu Pro 535 Pro	Cys 440 Gly His Leu Ile Arg 520 Ser	ASP Glu Tyr Lys 505 Pro Ala Pro	Arg Glu Pro Asn 490 Ala Ala Ser Val	Leu Glu 475 Ile Asn Val Ser Thr	Arg Val 460 Asn Gly Cys Thr Gly 540 Thr	Lys 445 Phe Leu Asp Pro Asn 525 Asn	A30 Ser Ser Met Cys 510 Gly Glu	Arg Trp Pro Val 495 Ile Met Thr	Pro Leu Thr 480 His Ser Ser Thr Asp 560
Arg Lys 465 Gln Ala Arg Gln Phe 545	Phe Ser 450 Cys Ile Ala Gln Leu 530 Ser	Gly 435 Glu Ala Ile Arg Asn 515 Pro	420 Val Thr Lys Pro Gly 500 Lys Ser	Glu Gly Gly 485 Lys Ser Ile Gly	Leu Glu Gln 470 Thr Trp Val Asn	Asp Met 455 Ser Ala Gly Leu Pro 535 Pro	Cys 440 Gly His Leu Ile Arg 520 Ser	ASP Glu Tyr Lys 505 Pro Ala Pro	Arg Glu Pro Asn 490 Ala Ala Ser Val	Glu 475 Ile Asn Val Ser Thr 555 Arg	Arg Val 460 Asn Gly Cys Thr Gly 540 Thr	Lys 445 Phe Leu Asp Pro Asn 525 Asn	A30 Ser Ser Met Cys 510 Gly Glu	Arg Trp Pro Val 495 Ile Met Thr	Pro Leu Thr 480 His Ser Ser Thr
Arg Lys 465 Gln Ala Arg Gln Phe 545	Phe Ser 450 Cys Ile Ala Gln Leu 530 Ser Val	Gly 435 Glu Ala Ile Arg Asn 515 Pro Gly	420 Val Thr Lys Pro Glys 500 Lys Ser Gly Lys	Cys Glu Gly Gly 485 Lys Ser Ile Gly Ala 565	Leu Glu Gln 470 Thr Val Asn Gly 550 Asp	Asp Met 455 Ser Ala Gly Leu Pro 535 Pro	Cys 440 Gly His Leu Ile Arg 520 Ser Ala	425 Tyr Asp Glu Tyr Lys 505 Pro Ala Pro	Arg Glu Pro Asn 490 Ala Ala Ser Val Ile 570	Glu Glu 475 Ile Asn Val Ser Thr 555 Arg	Arg Val 460 Asn Cys Thr Gly 540 Thr	Lys 445 Phe Leu Asp Pro Asn 525 Asn	430 Ser Ser Met Cys 510 Gly Glu Glu	Arg Trp Pro Val 495 Ile Met Thr Pro 575	Pro Leu Thr 480 His Ser Thr Asp 560 Leu
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Lys 465 Gln Ala Arg Gln Phe 545 His	Phe Ser 450 Cys Ile Ala Gln Leu 530 Ser Val	Gly 435 Glu Ala Ile Arg Asn 515 Pro Gly Pro	420 Val Thr Lys Pro Glyy 500 Lys Ser Gly Lys	Cys Glu Gly Gly 485 Lys Ser Ile Gly Ala 565 Ser	Leu Glu Gln 470 Thr Val Asn Gly 550 Asp	Asp Met 455 Ser Ala Gly Leu Pro 535 Pro Ser	Cys 440 Gly His Leu Ile Arg 520 Ser Ala Thr	425 Tyr Asp Glu Tyr Lys 505 Pro Ala Pro Asp	Arg Glu Pro Asn 490 Ala Ala Ser Val Ile 570 Asn	Glu Glu 475 Ile Asn Val Ser Thr 555 Arg	Arg Val 460 Asn Gly Cys Thr Gly 540 Thr	Lys 445 Phe Leu Asp Pro Asn 525 Asn Pro Glu	430 Ser Ser Met Cys 510 Gly Glu Glu Glu Lys 590	Arg Trp Pro Val 495 Ile Met Thr Pro 575 Ala	Pro Leu Thr 480 His Ser Thr Asp 560 Leu Ile
Arg Lys 465 Gln Ala Arg Gln Phe 545 His Lys	Phe Ser 450 Cys Ile Ala Gln Leu 530 Ser Val Thr	Gly 435 Glu Ala Ile Arg Asn 515 Pro Gly Pro Asp Pro 595	420 Val Thr Lys Pro Glyy 500 Lys Ser Gly Lys Ser Ser Cys	Cys Glu Gly 485 Lys Ser Ile Gly Alaa 565 Ser	Leu Glu Gln 470 Thr Trp Val Asn Gly 550 Asp Ala	Asp Met 455 Ser Ala Gly Leu Pro 535 Pro Ser Ser	Cys 440 Gly His Leu Ile Arg 520 Ser Ala Thr Asn Ala 600	425 Tyr Asp Glu Tyr Lys 505 Pro Ala Pro Asp Ser 585 Pro	Arg Glu Pro Asn 490 Ala Ser Val Ile 570 Asn Pro	Leu Glu Glu 475 Ile Asn Val Ser Thr 555 Arg Ser	Arg Val 460 Asn Gly Cys Thr Gly 540 Thr Ser Glu Ser	Lys 445 Phe Leu Asp Pro Asn 525 Asn Pro Glu Leu Ala 605	430 Ser Ser Met Cys 510 Glu Glu Glu Lys 590 Leu	Arg Trp Pro Val 495 Ile Met Thr Pro Pro 575 Ala His	Pro Leu Thr 480 His Ser Ser Thr Asp 560 Leu Ile
Arg Lys 465 Gln Ala Arg Gln Phe 545 His Lys	Phe Ser 450 Cys Ile Ala Gln Leu 530 Ser Val Thr	Gly 435 Glu Ala Ile Arg Asn 515 Pro Gly Pro Asp Pro 595	420 Val Thr Lys Pro Glyy 500 Lys Ser Gly Lys Ser Ser Cys	Cys Glu Gly 485 Lys Ser Ile Gly Alaa 565 Ser	Glu Gln 470 Thr Trp Val Asn Gly 550 Asp	Asp Met 455 Ser Ala Gly Leu Pro 535 Pro Ser Ser	Cys 440 Gly His Leu Ile Arg 520 Ser Ala Thr Asn Ala 600	425 Tyr Asp Glu Tyr Lys 505 Pro Ala Pro Asp Ser 585 Pro	Arg Glu Pro Asn 490 Ala Ser Val Ile 570 Asn Pro	Leu Glu Glu 475 Ile Asn Val Ser Thr 555 Arg Ser	Arg Val 460 Asn Gly Cys Thr Gly 540 Thr Ser Glu Ser	Lys 445 Phe Leu Asp Pro Asn 525 Asn Pro Glu Leu Ala 605	430 Ser Ser Met Cys 510 Glu Glu Glu Lys 590 Leu	Arg Trp Pro Val 495 Ile Met Thr Pro Pro 575 Ala His	Pro Leu Thr 480 His Ser Ser Thr Asp 560 Leu Ile

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615
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Gly Ser Leu Arg Ser Val Leu Asn Lys Glu Ser His Ser Pro Phe Gly
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                  630
Leu Asp Ser Phe Asn Ser Thr Ala Lys Val Ser Pro Leu Thr Pro Lys
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                                  650
Leu Phe Asn Ser Leu Leu Gly Pro Thr Ala Ser Asn Asn Lys Thr
                               665
Glu Gly Ser Ser Leu Arg Asp Leu Leu His Ser Gly Pro Gly Lys Leu
                           680
Pro Gln Thr Pro Leu Asp Thr Gly Ile Pro Phe Pro Pro Val Phe Ser
                       695
                                          700
Thr Ser Ser Ala Gly Val Lys Ser Lys Ala Ser Leu Pro Asn Phe Leu
                                      715
                   710
Asp His Ile Ile Ala Ser Val Val Glu Asn Lys Lys Thr Ser Asp Ala
                                   730
Ser Lys Arg Ala Cys Asn Leu Thr Asp Thr Gln Lys Glu Val Lys Glu
                              745
Met Val Met Gly Leu Asn Val Leu Asp Pro His Thr Ser His Ser Trp
                           760
Leu Cys Asp Gly Arg Leu Leu Cys Leu His Asp Pro Ser Asn Lys Asn
                       775
                                           780
Asn Trp Lys Ile Phe Arg Glu Cys Trp Lys Gln Gly Gln Pro Val Leu
                   790
                                       795
Val Ser Gly Val His Lys Lys Leu Lys Ser Glu Leu Trp Lys Pro Glu
                                   810
Ala Phe Ser Gln Glu Phe Gly Asp Gln Asp Val Asp Leu Val Asn Cys
                               825
Arg Asn Cys Ala Ile Ile Ser Asp Val Lys Val Arg Asp Phe Trp Asp
                          840
Gly Phe Glu Ile Ile Cys Lys Arg Leu Arg Ser Glu Asp Gly Gln Pro
                       855
Met Val Leu Lys Leu Lys Asp Trp Pro Pro Gly Glu Asp Phe Arg Asp
                                       875
Met Met Pro Thr Arg Phe Glu Asp Leu Met Glu Asn Leu Pro Leu Pro
                                   890
Glu Tyr Thr Lys Arg Asp Gly Arg Leu Asn Leu Ala Ser Arg Leu Pro
                               905
Ser Tyr Phe Val Arg Pro Asp Leu Gly Pro Lys Met Tyr Asn Ala Tyr
Gly Leu Ile Thr Ala Glu Asp Arg Arg Val Gly Thr Thr Asn Leu His
                       935
Leu Asp Val Ser Asp Ala Val Asn Val Met Val Tyr Val Gly Ile Pro
                                       955
                   950
Ile Gly Glu Gly Ala His Asp Glu Glu Val Leu Lys Thr Ile Asp Glu
                                   970
                965
Gly Asp Ala Asp Glu Val Thr Lys Gln Arg Ile His Asp Gly Lys Glu
                               985
Lys Pro Gly Ala Leu Trp His Ile Tyr Ala Ala Lys Asp Ala Glu Lys
                           1000
Ile Arg Glu Leu Leu Arg Lys Val Gly Glu Glu Gln Gly Gln Glu Asn
                                          1020
                       1015
Pro Pro Asp His Asp Pro Ile His Asp Gln Ser Trp Tyr Leu Asp Gln
                                       1035
                   1030
Thr Leu Arg Lys Arg Leu Tyr Glu Glu Tyr Gly Val Gln Gly Trp Ala
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1050
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Ile Val Gln Phe Leu Gly Asp Ala Val Phe Ile Pro Ala Gly Ala Pro
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            1060
His Gln Val His Asn Leu Tyr Ser Cys Ile Lys Val Ala Glu Asp Phe
        1075
                            1080
                                                1085
Val Ser Pro Glu His Val Lys His Cys Phe Arg Leu Thr Gln Glu Phe
                                            1100
                        1095
Arg His Leu Ser Asn Thr His Thr Asn His Glu Asp Lys Leu Gln Val
                                        1115
Lys Asn Ile Ile Tyr His Ala Val Lys Asp Ala Val Gly Thr Leu Lys
                                                        1135
                1125
                                    1130
Ala His Glu Ser Lys Leu Ala Arg Ser
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                                1145
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actcatttgc cccgcaggta gatcttgggg gtctgccagc cttcgggggc ttcctttagc
cccgccttca gccagatgcg cctcaggtct ttctcgaact tgatctgctt gcgtctcagg
egiceeteet ggacetteet eegeaggaac egegiettet teaceagett eeggiacitg
tggtggttca tetteegeeg geggatette ageaegtttt tgeaetgaat ttgaggegea
teegegacge etteateece etgeteggee cetteeceta tetggetggg eggacaetgg
taggattgcg gtggagccac agtccctgcg gtcccggtat ccagtctggg caggaagcag
cgggccgtga gccagctctc cagggggctg acggacatct tcctggggac cagcatctcc
tecageteca getgggeece ettgegaggg agagaggeeg eeetacetgg geeggeegge
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geoccetocc egggttcacc cccgcgcgaa tcgcgttgcc tggcgcccgg accetctcgg
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<210> 4596

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<213> Homo sapiens
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Arg Lys Gln Arg Ala Val Ser Gln Leu Ser Arg Gly Leu Thr Asp Ile
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                                                     30
Phe Leu Gly Thr Ser Ile Ser Ser Ser Ser Ser Trp Ala Pro Leu Arg
        35
                            40
Gly Arg Glu Ala Ala Leu Pro Gly Pro Ala Gly Asp Xaa Ala Val Lys
                        55
                                            60
Gly Pro Ala Asp Pro Ala Ala Gln His Ser Arg Asp Gly Gln Gly Gly
                                        75
                                                             RΛ
                    70
Trp Pro Pro Ala Gln Gly Thr Ala Ser Thr Ala Gly Lys Ser Gly Ala
                85
                                    90
Pro Gly Ala Trp Ser Val Gly Gly Ala Thr Gly Pro Arg Gly Ala Lys
                                105
                                                     110
Gly Pro Arg Thr Gly Arg Pro Ala Pro Ser Pro Gly Ser Pro Pro Arg
                            120
Glu Ser Arg Cys Leu Ala Pro Gly Pro Ser Arg Leu Asp Pro Gly Pro
                        135
                                            140
Ala Xaa Ala Ala Ala Pro Gly Ala Leu Arg Pro Pro Ala Asp Pro Ser
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                                        155
                                                            160
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Gln Ala Arg Pro Arg Arg Gly Ser Asn
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<210> 4597
<211> 515
<212> DNA
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qqqacactca tgctcagtga ctgatgggat ggggggtaca aagtcccagc cacgtgattc
tgggaggcca ttccagctca caactcctgg gccctgggga gtcggccgtg ggacctgcct
cacageteag etecteetet eggeeceatt etgeeteete eeggeeettt eecaggeagt
aagcccaagg aactccttaa gaaacatcct cactctgaac tccactgcag agccttcttc
ctgggaaagc agggagcgcc ccctgcaatc acgtaatgtt tactcatccg cctccttctc
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<210> 4598

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<212> PRT
<213> Homo sapiens
<400> 4598
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Thr Lys Ser Gln Pro Arg Asp Ser Gly Arg Pro Phe Gln Leu Thr Thr
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                                                     30
            20
Pro Gly Pro Trp Gly Val Gly Arg Gly Thr Cys Leu Thr Ala Gln Leu
        35
                            40
Leu Leu Ser Ala Pro Phe Cys Leu Leu Pro Ala Leu Ser Gln Ala Val
                        55
Ser Pro Arg Asn Ser Leu Arg Asn Ile Leu Thr Leu Asn Ser Thr Ala
                                                             80
                    70
Glu Pro Ser Ser Trp Glu Ser Arg Glu Arg Pro Leu Gln Ser Arg Asn
Val Tyr Ser Ser Ala Ser Phe Ser Glu His Leu Asp Gly Gly Cys Ser
                                105
Pro Leu Val Leu Gln Ser Leu Ala Arg Arg Ile Ser Ser Thr Trp Leu
                                                 125
        115
                            120
Val Asp Gln Ser Leu Arg Glu
    130
                        135
<210> 4599
<211> 2314
<212> DNA
<213> Homo sapiens
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ggaggcaggt teegcacgaa ataaatcaga atgagttatg cagaaaaacc egatgaaatc
acgaaagatg agtggatgga aaagctcaat aacttgcatg tccagagagc agacatgaac
cgcctcatca tgaactacct ggtcacagag ggctttaagg aagcagcgga gaagtttcga
atggaatctg gaatcgaacc tagtgtggat ctggaaacac ttgatgaacg aatcaagatc
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 ccagagetet tggacacaaa eeggtatett taetteeatt tgcagcaaca geatttgate
 gagetgatee gecageggga gacagaggeg gegetggagt ttgcacagae teagetggeg
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 gcctttgaca gtcccgagga gtcgcccttc ggagacctcc tccacaccat gcagaggcag
 aaggtgtgga gtgaagttaa ccaagctgtg ctagattatg aaaatcgcga gtcaacaccc
 720
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aaactqqcaa aattactgaa actactactt tgggctcaga acgagctgga ccagaagaaa
gtaaaatatc ccaaaatgac agacctcagc aagggtgtga ttgaggagcc caagtagcgc
840
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gacctggcat ctttttatag ggaaaaatgg cctttgtagg cagtggaaaa cttgcaagga
aagctgccgt ctctttggca gtctgatgca gagcctgcac tctggcactc gctgaagaat
1080
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1140
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1200
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cttacccctg tggtttttgt gtttttttt ttttcttttt ccataggaaa gaatatataa
1320
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1740
gcagagcccc agcaggtggt gcacgactgt tggcggaagg aacgcgtgtt catcctcagt
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gatotgocot coagoatoto ggoagoatot catootocat cgtcageogg ctotgoogat
1860
qtcctqcttc tgttcactca cagaactgtc ccctgctccg tggtgggcag gagggaagtg
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gtgcagggct gcgtgcattg cctgcgagtc gggacagttg atgggcacat ggccttgtag
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2314
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<400> 4600
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            20
Ile Met Asn Tyr Leu Val Thr Glu Gly Phe Lys Glu Ala Ala Glu Lys
                                                45
                            40
Phe Arg Met Glu Ser Gly Ile Glu Pro Ser Val Asp Leu Glu Thr Leu
                        55
Asp Glu Arg Ile Lys Ile Arg Glu Met Ile Leu Lys Gly Gln Ile Gln
                    70
Glu Ala Ile Ala Leu Ile Asn Ser Leu His Pro Glu Leu Leu Asp Thr
                                    90
Asn Arg Tyr Leu Tyr Phe His Leu Gln Gln His Leu Ile Glu Leu
                                                    110
                                105
Ile Arg Gln Arg Glu Thr Glu Ala Ala Leu Glu Phe Ala Gln Thr Gln
                            120
Leu Ala Glu Gln Gly Glu Glu Ser Arg Glu Cys Leu Thr Glu Met Glu
                        135
Arg Thr Leu Ala Leu Leu Ala Phe Asp Ser Pro Glu Glu Ser Pro Phe
                                        155
                    150
145
Gly Asp Leu Leu His Thr Met Gln Arg Gln Lys Val Trp Ser Glu Val
                                     170
Asn Gln Ala Val Leu Asp Tyr Glu Asn Arg Glu Ser Thr Pro Lys Leu
                                                     190
                                 185
Ala Lys Leu Leu Lys Leu Leu Trp Ala Gln Asn Glu Leu Asp Gln
                             200
        195
 Lys Lys Val Lys Tyr Pro Lys Met Thr Asp Leu Ser Lys Gly Val Ile
                         215
 Glu Glu Pro Lys
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 <212> DNA
 <213> Homo sapiens
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 tttctaaaga gtgaccgagt ggccagaatg gtacagagtg gagggtgttc tgctaatgac
 ttcagagaag tatttaagaa aaacatagaa aaacgtgtgc ggagtttgcc agaaatagat
 ggcttgagca aagagacagt gttgagctca tggatagcca aatatgatgc catttacaga
 300
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ggtgaagagg acttgtgcaa acagccaaat agaatggccc taagtgcagt gtctgaactt
attotgagoa aggaacaact otatgaaatg tttcagcaga ttotgggtat taaaaaacta
gaacaccage teetttataa tgeatgteag etggataaeg eagatgaaca ageageeeag
atcagaaggg aacttgatgg ccggctgcaa ttggcagata aaatggcaaa ggaaagaaaa
ttccccaaat ttatagcaaa agatatggag aatatgtata tagaagagtt gcggtcttca
gtgaatttgc taatggccaa tttggaaagt cttccagttt cgaaaggtgg tccggaattt
aaattacaaa aattaaaacg ttcacagaac tctgcatttt tggacatagg agatgagaat
gagattcagc tgtcaaagtc cgacgtggta ctgtcattca ccttagagat tgtcataatg
gaagtgcaag geetgaagte agttgeteee aategaattg tttactgtae aatggaagtg
gaaggagaaa aacttcagac agaccaggcc gaagcctcaa ggccacaatg gggggactca
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<212> PRT
<213> Homo sapiens
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Phe Leu Asn Gly Glu Thr Gln Ile Val Ala Asp Glu Ala Phe Cys Asn
                                 25
Ala Val Arg Ser Tyr Tyr Glu Val Phe Leu Lys Ser Asp Arg Val Ala
                             40
Arg Met Val Gln Ser Gly Gly Cys Ser Ala Asn Asp Phe Arg Glu Val
                                             60
Phe Lys Lys Asn Ile Glu Lys Arg Val Arg Ser Leu Pro Glu Ile Asp
                                         75
                     70
Gly Leu Ser Lys Glu Thr Val Leu Ser Ser Trp Ile Ala Lys Tyr Asp
                                     90
                 85
 Ala Ile Tyr Arg Gly Glu Glu Asp Leu Cys Lys Gln Pro Asn Arg Met
                                 105
             100
 Ala Leu Ser Ala Val Ser Glu Leu Ile Leu Ser Lys Glu Gln Leu Tyr
                                                 125
 Glu Met Phe Gln Gln Ile Leu Gly Ile Lys Lys Leu Glu His Gln Leu
                                             140
                         135
 Leu Tyr Asn Ala Cys Gln Leu Asp Asn Ala Asp Glu Gln Ala Ala Gln
                                         155
                     150
 Ile Arg Arg Glu Leu Asp Gly Arg Leu Gln Leu Ala Asp Lys Met Ala
                                     170
                 165
 Lys Glu Arg Lys Phe Pro Lys Phe Ile Ala Lys Asp Met Glu Asn Met
                                                     190
                                 185
 Tyr Ile Glu Glu Leu Arg Ser Ser Val Asn Leu Leu Met Ala Asn Leu
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205
        195
                            200
Glu Ser Leu Pro Val Ser Lys Gly Gly Pro Glu Phe Lys Leu Gln Lys
                                            220
Leu Lys Arg Ser Gln Asn Ser Ala Phe Leu Asp Ile Gly Asp Glu Asn
                                        235
                    230
225
Glu Ile Gln Leu Ser Lys Ser Asp Val Val Leu Ser Phe Thr Leu Glu
                245
Ile Val Ile Met Glu Val Gln Gly Leu Lys Ser Val Ala Pro Asn Arg
                                265
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Arg Gln Arg Asn Arg Leu Arg Leu Glu Glu Asp Lys Pro Ala Val Glu
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 Arg Cys Leu Glu Glu Leu Val Phe Gly Asp Val Glu Asn Asp Glu Asp
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 Ser Gly Asp Ser Glu Val Glu Asn Glu Ala Lys Gly Asn Phe Pro Pro
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 Gln Lys Lys Pro Val Trp Val Asp Glu Glu Asp Glu Asp Glu Glu Met
                             120
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 Val Asp Met Met Asn Asn Arg Phe Arg Lys Asp Met Met Lys Asn Ala
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 Ser Glu Ser Lys Leu Ser Lys Asp Asn Leu Lys Lys Arg Leu Lys Glu
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 Glu Phe Gln His Ala Met Gly Gly Val Pro Ala Trp Ala Glu Thr Thr
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Lys Arg Lys Thr Ser Ser Asp Asp Glu Ser Glu Glu Asp Glu Asp Asp
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Pro Thr Val Ala Arg Ile Ser Ser Val Gln Phe His Pro Gly Ala Gln
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                                       235
Ile Val Met Val Ala Gly Leu Asp Asn Ala Val Ser Leu Phe Gln Val
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                                  250
Asp Gly Lys Thr Asn Pro Lys Ile Gln Ser Ile Tyr Leu Glu Arg Phe
                              265
Pro Ile Phe Lys Ala Cys Phe Ser Ala Asn Gly Glu Glu Val Leu Ala
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Thr Ser Thr His Ser Lys Val Leu Tyr Val Tyr Asp Met Leu Ala Gly
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Lys Leu Ile Pro Val His Gln Val Arg Gly Leu Lys Glu Lys Ile Val
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Arg Ser Phe Glu Val Ser Pro Asp Gly Ser Phe Leu Leu Ile Asn Gly
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Ile Ala Gly Tyr Leu His Leu Leu Ala Met Lys Thr Lys Glu Leu Ile
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Gly Ser Met Lys Ile Asn Gly Arg Val Ala Ala Ser Thr Phe Ser Ser
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Asp Ser Lys Lys Val Tyr Ala Ser Ser Gly Asp Gly Glu Val Tyr Val
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Ser Leu Tyr Gly Leu Ser Ile Ala Thr Ser Arg Asn Gly Gln Tyr Val
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Ala Cys Gly Ser Asn Cys Gly Val Val Asn Ile Tyr Asn Gln Asp Ser
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Cvs Leu Gln Glu Thr Asn Pro Lys Pro Ile Lys Ala Ile Met Asn Leu
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                                              445
Val Thr Gly Val Thr Ser Leu Thr Phe Asn Pro Thr Thr Glu Ile Leu
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                                          460
Ala Ile Ala Ser Glu Lys Met Lys Glu Ala Val Arg Leu Val His Leu
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Pro Ser Cys Thr Val Phe Ser Asn Phe Pro Val Ile Lys Asn Lys Asn
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Ile Ser His Val His Thr Met Asp Phe Ser Pro Arg Ser Gly Tyr Phe
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Asp Phe Leu Ile Phe Thr Thr Gln Ile Leu Thr Ile Leu Gln Leu Arg
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Ser Leu Asn Ile Ile Tyr Asn Lys Gln Asn Leu Val Asn Leu Gln Lys
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Ser Asn Ala Leu Lys Lys His Gln Ser Leu Cys Met Cys Arg Thr Asp
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Ser Val Ser Leu Leu
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240
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Arg Lys Asp Met Asp Glu Val Glu Glu Lys Ser Lys Asp Val Ile Asn
Phe Thr Ala Glu Lys Leu Ser Val Asp Glu Val Ser Gln Leu Val Ile
                        55
Ser Pro Leu Cys Gly Ala Ile Ser Leu Phe Val Gly Thr Thr Arg Asn
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Asn Phe Glu Gly Lys Lys Val Ile Ser Leu Glu Tyr Glu Ala Tyr Leu
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Pro Met Ala Glu Asn Glu Val Arg Lys Ile Cys Ser Asp Ile Arg Gln
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Lys Trp Pro Val Lys His Ile Ala Val Phe His Leu Leu Gly Leu Val
                            120
Pro Val Ser Glu Ala Ser Thr Val Ile Ala Val Ser Ser Ala His Arg
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Ala Ala Ser Leu Glu Ala Val Ser Tyr Ala Ile Asp Ser Leu Lys Ala
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Lys Val Pro Ile Trp Lys Lys Glu Ile Tyr Glu Glu Ser Ser Thr Trp
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Gln Pro Thr Ala Glu Pro Gly Leu Gly Ala Val Val Arg Ser Ile Lys
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Val Ser Gly Tyr Leu Asn Leu Leu Ala Asn Thr Ile Asp Asn Phe Thr
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His Gly Leu Ala Val Ala Ala Ser Phe Leu Val Ser Lys Lys Ile Gly
                                         75
Leu Leu Thr Thr Met Ala Ile Leu Leu His Glu Ile Pro His Glu Val
                                     90
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Gly Asp Phe Ala Ile Leu Leu Arg Ala Gly Phe Asp Arg Trp Ser Ala
            100
                                 105
Ala Lys Leu Gln Leu Ser Thr Ala Leu Gly Gly Leu Leu Gly Ala Gly
                                                 125
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Phe Ala Ile Cys Thr Gln Ser Pro Lys Gly Val Glu Glu Thr Ala Ala
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                                             140
Trp Val Leu Pro Phe Thr Ser Gly Gly Phe Leu Tyr Ile Ala Leu Val
                                         155
                    150
Asn Val Leu Pro Asp Leu Leu Glu Glu Glu Asp Pro Trp Arg Ser Leu
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Gln Gln Leu Leu Leu Cys Ala Gly Ile Val Val Met Val Leu Phe
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Ser Leu Phe Val Asp
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<213> Homo sapiens

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Trp Glu Lys Leu Thr Asn Glu Ser Ser Trp Gln Pro Pro Gln Ala Pro
Pro Asp Trp Ala Ser Trp Leu Cys Cys Gln Asp Tyr Asp Pro Leu Pro
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Glu Ser Arg Arg Ser Pro Gln Ala Glu Arg Tyr Arg His Leu Cys Pro
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Tyr Leu Asn Gln Glu Val Pro
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Ile Gly Lys Lys Gly Glu Thr Val Lys Arg Ile Arg Glu Gln Ser Ser
                           40
Ala Arg Ile Thr Ile Ser Glu Gly Ser Cys Pro Glu Arg Ile Thr Thr
                        55
Ile Thr Gly Ser Thr Ala Ala Val Phe His Ala Val Ser Met Ile Ala
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Phe Lys Leu Asp Glu Asp Leu Cys Ala Ala Pro Ala Asn Gly Gly Asn
                                    90
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Val Ser Arg Pro Pro Val Thr Leu Arg Leu Val Ile Pro Ala Ser Gln
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Cys Gly Ser Leu Ile Gly Lys Ala Gly Thr Lys Ile Lys Glu Ile Arg
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Glu Thr Thr Gly Ala Gln Val Gln Val Ala Gly Asp Leu Leu Pro Asn
                       135
                                           140
Ser Thr Glu Arg Ala Val Thr Val Ser Gly Val Pro Asp Ala Ile Ile
                                        155
                    150
Leu Cys Val Arg Gln Ile Cys Ala Val Ile Leu Glu Ser Pro Pro Lys
                                   170
Gly Ala Thr Ile Pro Tyr His Pro Ser Leu Ser Leu Gly Thr Val Leu
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Leu Ser Ala Asn Gln Gly Phe Ser Val Gln Gly Gln Tyr Gly Ala Val
                           200
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Thr Pro Ala Glu Val Thr Lys Leu Gln Gln Leu Ser Ser His Ala Val
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Pro Phe Ala Thr Pro Ser Val Val Pro Gly Leu Asp Pro Gly Thr Gln
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Thr Ser Ser Gln Glu Phe Leu Val Pro Asn Asp Leu Ile Gly Cys Val
                                    250
Ile Gly Arg Gln Gly Ser Lys Ile Ser Glu Ile Arg Gln Met Ser Gly
                               265
Ala His Ile Lys Ile Gly Asn Gln Ala Glu Gly Ala Gly Glu Arg His
                                                285
                            280
Val Thr Ile Thr Gly Ser Pro Val Ser Ile Ala Leu Ala Gln Tyr Leu
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Ile Thr Ala Cys Leu Glu Thr Ala Lys Ser Thr Ser Gly Gly Thr Pro
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Gly Ser Ala Pro Ala Asp Leu Pro Thr Pro Phe Ser Pro Pro Leu Thr
                325
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Ala Leu Pro Thr Ala Pro Pro Gly Leu Leu Gly Thr Pro Tyr Ala Ile
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Ser Leu Ser Asn Phe Ile Gly Leu Lys Pro Val Pro Phe Leu Ala Leu
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cccccct

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365
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Pro Pro Ala Ser Pro Gly Pro Pro Pro Gly Leu Ala Ala Tyr Thr Ala
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1200
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aacatggccc tcggcaagaa ggcggctgac agcctgcagc agaatctgca gcgggactac
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Asp Pro Trp Lys Glu Glu Thr Asp Thr Asp Leu Glu Val Val Leu Glu
                            40
Lys Lys Gly Asn Met Asp Glu Ala His Ile Asp Gln Val Arg Arg Lys
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Ala Leu Gln Glu Glu Ile Asp Arg Glu Ser Gly Lys Thr Glu Ala Ser
                    70
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65
Glu Thr Arg Lys Trp Thr Gly Thr Gln Phe Gly Gln Trp Asp Thr Ala
Gly Phe Glu Asn Glu Asp Gln Lys Leu Lys Phe Leu Arg Leu Met Gly
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100
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Gly Phe Lys Asn Leu Ser Pro Ser Phe Ser Arg Pro Ala Ser Thr Ile
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Ala Arg Pro Asn Met Ala Leu Gly Lys Lys Ala Ala Asp Ser Leu Gln
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                        135
Gln Asn Leu Gln Arg Asp Tyr Asp Arg Ala Met Ser Trp Lys Tyr Ser
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                                        155
Arg Gly Ala Gly Leu Gly Phe Ser Thr Ala Pro Asn Lys Ile Phe Tyr
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Ile Asp Arg Asn Ala Ser Lys Ser Val Lys Leu Glu Asp
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Asp Met Gln Ala Leu Arg Arg Glu Glu Glu Arg Arg Gln Ala Glu Arg
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Glu Gln Glu Tyr Lys Arg Lys Gln Leu Glu Glu Gln Arg Gln Ser Glu
Arg Leu Gln Arg Gln Leu Gln Gln Glu His Ala Tyr Leu Lys Ser Leu
 Gln Gln Gln Gln Gln Gln Gln Leu Gln Lys Gln Gln Gln Gln
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 Leu Leu Pro Gly Asp Arg Lys Pro Leu Tyr His Tyr Gly Arg Gly Met
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<212> DNA <400> 4627

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<213> Homo sapiens

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Pro Glu Ala Lys Gln Glu Ile Leu Glu Asn Lys Asp Val Val Val Gln
                          40
His Val His Phe Asp Gly Leu Gly Arg Thr Lys Asp Asp Ile Ile Ile
                                         60
Cys Glu Ile Gly Asp Val Phe Lys Ala Lys Asn Leu Ile Glu Val Met
Arg Lys Ser His Glu Ala Arg Glu Lys Leu Leu Arg Leu Gly Ile Phe
Arg Gln Val Asp Val Leu Ile Asp Thr Cys Gln Gly Asp Gly Ala Leu
                             105
Pro Asn Gly Leu Asp Val Thr Phe Glu Val Thr Glu Leu Arg Arg Leu
       115
                          120
                                             125
Thr Gly Ser Tyr Asn Thr Met Val Gly Asn Asn Glu Gly Ser Met Val
                      135
Leu Gly Leu Lys Leu Pro Asn Leu Leu Gly Arg Ala Glu Lys Val Thr
                                     155
                  150
Phe Gln Phe Ser Tyr Gly Thr Lys Glu Thr Ser Tyr Gly Leu Ser Phe
                                 170
               165
Phe Lys Pro Arg Pro Gly Asn Phe Glu Arg Asn Phe Ser Val Asn Leu
                             185
Tyr Lys Val Thr Gly Gln Phe Pro Trp Ser Ser Leu Arg Glu Thr Asp
                          200
Arg Gly Met Ser Ala Glu Tyr Ser Phe Pro Ile Trp Lys Thr Ser His
                      215
                                         220
Thr Val Lys Trp Glu Gly Val Trp Arg Glu Leu Gly Cys Leu Ser Arg
                  230
                                     235
Thr Ala Ser Phe Ala Val Arg Lys Glu Ser Gly His Ser Leu Lys Ser
               245
                                 250
Ser Leu Ser His Ala Met Val Ile Asp Ser Arg Asn Ser Ser Ile Leu
                              265
Pro Arg Arg Gly Ala Leu Leu Lys Val Asn Gln Glu Leu Ala Gly Tyr
                          280
Thr Gly Gly Asp Val Ser Phe Ile Lys Glu Asp Phe Glu Leu Gln Leu
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                                         300
Asn Lys Gln Leu Ile Phe Asp Ser Val Phe Ser Ala Ser Phe Trp Gly
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Gly Met Leu Val Pro Ile Gly Asp Lys Pro Ser Ser Ile Ala Asp Arg
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Phe Tyr Leu Gly Gly Pro Thr Ser Val Arg Gly Phe Ser Met His Ser
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                                345
                                                    350
Ile Gly Pro Gln Ser Glu Gly Asp Tyr Leu Gly Gly Glu Ala Tyr Trp
                            360
                                                365
Ala Gly Gly Leu His Leu Tyr Thr Pro Leu Pro Phe Arg Pro Gly Gln
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Gly Gly Phe Gly Glu Leu Phe Arg Thr His Phe Phe Leu Asn Ala Gly
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Asn Leu Cys Asn Leu Asn Tyr Gly Glu Gly Pro Lys Ala His Ile Arg
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Lys Leu Ala Glu Cys Ile Arg Trp Ser Tyr Gly Ala Gly Ile Val Leu
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Arg Leu Gly Asn Ile Ala Arg Leu Glu Leu Asn Tyr Cys Val Pro Met
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Gly Val Gln Thr Gly Asp Arg Ile Cys Asp Gly Val Gln Phe Gly Ala
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Gly Ile Arg Phe Leu
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ageaccegca ttgcttcggc cctagtgcag gggcagcacg tgcgcactga accetggagt
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Arg Asp Gln Gly Ala Leu Ser Leu Ser Arg Met Gly Arg Asp Ala Ser
Ser Trp Ala Leu Arg Val Ser Val Phe Pro Gln Ile Gly Lys Met Arg
Gly Arg Gly Gly Tyr Trp Gly Gln Ala Ser Ala Gln Pro Trp Val Leu
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Leu Glu Pro Gly Leu Glu Pro Glu Val Gly Arg Val Ser Lys Leu Ser
                85
                                    90
Ser Trp Ile Pro Ile Cys Arg Thr Ala Pro Arg Thr Arg Ser Gly Val
            100
Arg Ala His Pro Leu Ala Arg Ile Leu Gly Ser Leu Gly His Lys Ala
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Gly Gln Gly Thr Arg Asp Pro Pro Thr Gln Glu Thr
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 720
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Asp Leu Gln Ile Ala Leu Ala Ser Phe Tyr Glu Asp Gly Gly Asp Glu
                             40
Asp Ile Val Thr Ile Ser Gln Ala Thr Pro Ser Ser Val Ser Arg Gly
Thr Ala Pro Ser Asp Asn Arg Val Thr Ser Phe Arg Asp Leu Ile His
                                         75
                    70
Asp Gln Asp Glu Asp Glu Glu Glu Glu Glu Gly Gln Arg Ser Arg Phe
                                     90
Tyr Ala Gly Gly Ser Glu Arg Ser Gly Gln Gln Ile Val Gly Pro Pro
            100
                                 105
Arg Lys Lys Ser Pro Asn Glu Leu Val Asp Asp Leu Phe Lys Gly Ala
                             120
Lys Glu His Gly Ala Val Ala Val Glu Arg Val Thr Lys Ser Pro Gly
                        135
                                             140
Glu Thr Ser Lys Pro Arg Pro Phe Ala Gly Gly Gly Tyr Arg Leu Gly
                                         155
                     150
Ala Ala Pro Glu Glu Glu Ser Ala Tyr Val Ala Gly Glu Lys Arg Gln
                                     170
                 165
His Ser Ser Gln Asp Val His Val Val Leu Lys Leu Trp Lys Ser Gly
                                                     190
                                 185
Phe Ser Leu Asp Asn Gly Glu Leu Arg Ser Tyr Gln Asp Pro Ser Asn
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                             200
Ala Gln Phe Leu Glu Ser Ile Arg Arg Gly Glu Val Pro Ala Glu Leu
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                         215
Arg Arg Leu Ala His Gly Gly Gln Val Asn Leu Asp Met Glu Asp His
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Arg Asp Glu Asp Phe Val Lys Pro Lys Gly Ala Phe Lys Ala Phe Thr
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Gly Glu Gly Gln Lys Leu Gly Ser Thr Ala Pro Gln Val Leu Ser Thr
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265
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Ser Ser Pro Ala Gln Gln Ala Glu Asn Glu Ala Lys Ala Ser Ser Ser
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Ile Leu Ile Asp Glu Ser Glu Pro Thr Thr Asn Ile Gln Ile Arg Leu
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                                            300
Ala Asp Gly Gly Arg Leu Val Gln Lys Phe Asn His Ser His Arg Ile
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                                                             320
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Ser Asp Ile Arg Leu Phe Ile Val Asp Ala Arg Pro Ala Met Ala Ala
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                                    330
Thr Ser Phe Ile Leu Met Thr Thr Phe Pro Asn Lys Glu Leu Ala Asp
                                345
                                                     350
            340
Glu Ser Gln Thr Leu Lys Glu Ala Asn Leu Leu Asn Ala Val Ile Val
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Gln Arg Leu Thr
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<211> 242

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Arg Ile Cys Ile Gln Ala Ile Leu Gln Asp Lys Pro Lys Ile Ala Thr
                                25
Ala Asn Leu Gly Lys Phe Leu Glu Leu Leu Arg Ser His Gln Ser Arg
                            40
Pro Ala Lys Cys Leu Thr Ile Met Trp Ala Leu Gly Gln Ala Gly Phe
                        55
Ala Asn Leu Thr Glu Gly Leu Lys Val Trp Leu Gly Ile Met Leu Pro
                    70
                                        75
Val Leu Gly Ile Lys Ser Leu Ser Pro Phe Ala Ile Thr Tyr Leu Asp
                                    90
                85
Arg Leu Leu Met His Pro Asn Leu Thr Lys Gly Phe Gly Met Ile
            100
                                 105
Gly Pro Lys Asp Phe Phe Pro Leu Leu Asp Phe Ala Tyr Met Pro Asn
                                                 125
                            120
Asn Ser Leu Thr Pro Ser Leu Gln Glu Gln Leu Cys Gln Leu Tyr Pro
                        135
                                             140
Arg Leu Lys Val Leu Ala Phe Gly Ala Lys Pro Asp Ser Thr Leu His
                    150
                                        155
Thr Tyr Phe Pro Ser Phe Leu Ser Arg Ala Thr Pro Ser Cys Pro Pro
                165
                                     170
Glu Met Lys Lys Glu Leu Leu Ser Ser Leu Thr Glu Cys Leu Thr Val
                                                     190
                                185
            180
Asp Pro Leu Ser Ala Ser Val Trp Arg Gln Leu Tyr Pro Lys His Leu
                                                 205
        195
                            200
Ser Gln Ser Ser Leu Leu Leu Glu His Leu Leu Ser Ser Trp Glu Gln
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                                             220
 Ile Pro Lys Lys Val Gln Lys Ser Leu Gln Glu Thr Ile Gln Ser Leu
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Lys Leu
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 300
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gectaccetg gecacagtea gtteccatte teatttteta agaattttat cacaaaacag
360
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384
<210> 4636
<211> 108
<212> PRT
<213> Homo sapiens
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Met Leu Gly Gly Pro Val Cys Ser Tyr Glu Leu Gly Gly Cys Pro Val
Thr Arg Val Leu Gly Gln Pro Arg Lys Leu Phe Ser Ile Gly Trp Gly
Lys Glu Val Lys Trp Gly Pro Arg Arg Lys Ala Gly Gly Val Trp Ala
        35
Glu Pro Ala Ser Gly Gly Leu Pro Pro Pro Glu Asp Glu Phe Cys Ser
Pro Gly Val Cys Thr Leu Thr Leu Ala His Ser Leu Thr His Lys Thr
                                        75
Leu Thr Leu Cys Phe Phe Trp Gly Glu Gly Gly His Trp Gln Lys Arg
                85
Leu Pro Trp Pro Gln Ser Val Pro Ile Leu Ile Phe
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<213> Homo sapiens
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Glu Asp Gly Gly Ala Leu Arg Gly Glu Val Ile Pro Glu His Glu Phe
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Leu Arg Arg Ser Phe Ala Leu Val Ala Gln Ala Arg Val Gln Trp Arg
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Asp Leu Ser Ser Leu Gln Pro Pro Pro Pro Arg Leu Lys Arg Phe Ser
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His Leu Ser Leu Pro Ser Ser
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Gln Trp Asn Tyr Cys Thr Leu Ser Gln Glu Ile Leu Arg Arg Pro Ile
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Val Ala Cys Glu Leu Gly Arg Leu Tyr Asn Lys Asp Ala Val Ile Glu
 Phe Leu Leu Asp Lys Ser Ala Glu Lys Ala Leu Gly Lys Ala Ala Ser
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 His Ile Lys Ser Ile Lys Asn Val Thr Glu Leu Lys Leu Ser Asp Asn
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 Pro Ala Trp Glu Gly Asp Lys Gly Asn Thr Lys Gly Asp Lys His Asp
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 Asp Leu Gln Arg Ala Arg Phe Ile Cys Pro Val Val Gly Leu Glu Met
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Gly Ala Ala Phe Gln Glu Asp Asp Val Ile Met Leu Asn Gly Thr Lys
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Glu Asp Val Asp Val Leu Lys Thr Arg Met Glu Glu Arg Arg Leu Arg
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                                185
Ala Lys Leu Glu Lys Lys Thr Lys Lys Pro Lys Ala Ala Glu Ser Val
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Ser Lys Pro Asp Val Ser Glu Glu Ala Pro Gly Pro Ser Lys Val Lys
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                                             220
Thr Gly Lys Pro Glu Glu Ala Ser Leu Asp Ser Arg Glu Lys Lys Thr
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Asn Leu Ala Pro Lys Ser Thr Ala Met Asn Glu Ser Ser Ser Gly Lys
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Ala Gly Lys Pro Pro Cys Gly Ala Thr Lys Arg Ser Ile Ala Asp Ser
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Glu Glu Ser Glu Ala Tyr Lys Ser Leu Phe Thr Thr His Ser Ser Ala
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Gln Glu Asp Asp Met Lys Thr Leu Val Ser Glu Thr Ile Arg Arg Phe
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Gly Arg Leu Asp Cys Val Val Asn Asn Ala Gly His His Pro Pro Pro
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Gln Arg Pro Glu Glu Thr Ser Ala Gln Gly Phe Arg Gln Leu Leu Glu
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Leu Asn Leu Leu Gly Thr Tyr Thr Leu Thr Lys Leu Ala Leu Pro Tyr
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                           120
Leu Arg Lys Ser Gln Gly Asn Val Ile Asn Ile Ser Ser Leu Val Gly
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                                           140
Ala Ile Gly Gln Ala Gln Ala Val Pro Tyr Val Ala Thr Lys Gly Ala
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Val Thr Ala Met Thr Lys Ala Leu Ala Leu Asp Glu Ser Pro Tyr Gly
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Val Arg Val Asn Cys Ile Ser Pro Gly Asn Ile Trp Thr Pro Leu Trp
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                                                   190
            180
Glu Glu Leu Ala Ala Leu Met Pro Asp Pro Arg Ala Thr Ile Arg Glu
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                                               205
Gly Met Leu Ala Gln Pro Leu Gly Arg Met Gly Gln Pro Ala Glu Val
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                                           220
Gly Ala Ala Ala Val Phe Leu Ala Ser Glu Ala Asn Phe Cys Thr Gly
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                                       235
Ile Glu Leu Leu Val Thr Gly Gly Ala Glu Leu Gly Tyr Gly Cys Lys
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Ala Leu Gln Leu His Pro Asp Arg Asn Pro Asp Asp Pro Gln Ala Gln
Glu Lys Phe Gln Asp Leu Gly Ala Ala Tyr Glu Val Leu Ser Asp Ser
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Glu Lys Arg Lys Gln Tyr Asp Thr Tyr Gly Glu Glu Gly Leu Lys Asp
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Gly His Gln Ser Ser His Gly Asp Ile Phe Ser His Phe Phe Gly Asp
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Phe Gly Phe Met Phe Gly Gly Thr Pro Arg Gln Gln Asp Arg Asn Ile
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Pro Arg Gly Ser Asp Ile Ile Val Asp Leu Glu Val Thr Leu Glu Glu
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Val Tyr Ala Gly Asn Phe Val Glu Val Val Arg Asn Lys Pro Val Ala
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Arg Gln Ala Pro Gly Lys Arg Lys Cys Asn Cys Arg Gln Glu Met Arg
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Thr Thr Gln Leu Gly Pro Gly Arg Phe Gln Met Thr Gln Glu Val Val
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Cys Asp Glu Cys Pro Asn Val Lys Leu Val Asn Glu Glu Arg Thr Leu
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Glu Val Glu Ile Glu Pro Gly Val Arg Asp Gly Met Glu Tyr Pro Phe
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Phe Arg Ile Lys Val Val Lys His Pro Ile Phe Glu Arg Arg Gly Asp
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                                    250
Asp Leu Tyr Thr Asn Val Thr Ile Ser Leu Val Glu Ser Leu Val Gly
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 Phe Glu Met Asp Ile Thr His Leu Asp Gly His Lys Val His Ile Ser
                            280
 Arg Asp Lys Ile Thr Arg Pro Gly Ala Lys Leu Trp Lys Lys Gly Glu
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Gly Leu Pro Asn Phe Asp Asn Asn Ile Lys Gly Ser Leu Ile Ile
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Glu Gly Ile Lys Gln Leu Leu Lys Gln Gly Ser Val Gln Lys Val Tyr
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Glu Met Leu Leu Ser Arg Leu Ser Arg Tyr Arg Ala Ser Pro Ser Ala
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Thr Leu Ala Ala Leu Thr Gly Ser Thr Ile Ser Asn Thr Leu Lys Glu
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Asp Gln Ala Ala Asn Thr Ser Cys Gly Leu Pro Leu Lys Met Leu Arg
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Lys Thr Pro Ile Tyr Thr Cys Gly Thr Tyr Leu Val Met Leu Val Pro
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Pro Pro Gly Gly Ser Gly Ser Ser Ala Thr Arg Ser Leu Phe Gly Gly
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D	130	717	175.1	Uio	Ser		Thr	Tle	Phe	Phe		Ile	Ser	Asp	Leu
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145	7.00	т1 о	Cl.	Dro	Met	Tyr	Gln	Tvr	Ser		Thr	Trp	Phe	Ile	Asn
Ald	ASII	TTC	Giu	165	N.C.C	-1-		-1-	170			-		175	
	m		11 i a		Leu	Thr	Hie	Sor		Lvs	Ser	Glu	Glu		Asn
Leu	TYL	met	180	ser	пец	1111	1113	185		-,-			190		
_		-1.	180	m	Ile	т1 о	7 en		Dhe	Thr	Len	Ser		Tvr	Asn
Leu	Arg		гуу	IAT	116	116	200	1113	FIIC	****	LCu	205		-1-	
_		195			Leu	Dha		Tira	n an	Tire	T 411		Dhe	Ser	Len
Asn		Cys	Arg	ser	Leu		GIU	цуь	нар	цуз	220	LCu	1110	001	
	210					215	•	a1	·			T10	Thr	G1n	Glu.
	Leu	Thr	TTE	GIY	Ile	met	цуъ	GIII	цуз	235	GIU	110	****	oru	240
225					230				-1.				1 an	Dro	
Val	Trp	Tyr	Phe		Leu	Thr	GIY	GIY		мта	пеп	мар	Maii	255	LYL
				245		_			250		21-		212		TIO
Pro	Asn	Pro		Pro	Gln	Trp	Leu		GIU	гуз	ALA	пр	270	GIU	116
			260					265						774	T
Val	Arg		Ser	Ala	Leu	Pro		Leu	His	GIY	Leu		GIU	HIS	Leu
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Glu	Gln	Asn	Leu	Gly	Glu		Lys	Leu	He	Tyr	Asp	Ser	AIA	Trp	PFO
	290					295					300				
His	Glu	Glu	Gln	Leu	Pro	Gly	Ser	Trp	Lys		Ser	GIn	GIY	Leu	GIU
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Lys	Met	Val	Ile		Arg	Cys	Leu	Arg			Lys	Met	vai	Pro	Ala
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Val	Arg	Glu	Phe	Ile	Ala	Glu	His		Gly	Lys	Leu	Tyr	Ile	GIu	ALA
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Pro	Thr	Phe	Asp	Leu	Gln	Gly		Tyr	Asn	Asp	Ser		Cys	Cys	Ala
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Pro	Leu	Ile	Phe	Val	Leu	Ser	Pro	Ser	Ala	Asp	Pro	Met	Ala	GIA	Leu
	370					375					380			_	
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Ile	Ser	Leu	Gly	Gln	Gly	Gln	Gly	Pro	Ile	Ala	Ala	Lys	Met	Ile	Asn
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Asn	Ala	Ile	Lys	Asp	Gly	Thr	Trp	Val	Val	Leu	Gln	Asn	Cys	His	Leu
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Ala	Ala	Ser	Trp	Met	Pro	Thr	Leu	Glu	Lys	Ile	Cys			Val	Ile
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Val	Pro	Glu	Ser	Thr	Asn	Ala	Arg	Phe	Arg	Leu			Thr	Ser	Tyr
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Thr	Asn	Glu	Pro	Pro	Lys	Gly	Leu	Arg	Ala	Asn	Leu	Leu	Arg	Ser	Tyr
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545		. FIIC		. 020	550					555		-			560
Mot	Dhe	. T.o.	Asr	Asr	Tyr		Glu	Val	Pro			Ala	Let	Thr	Tyr
met	Pile	, net	. Asi		- 11	2,3	0								-

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Thr Leu Pro Arg Gln Ser Gly Gly Ser Gly Lys Ser Pro Gln Glu Val
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Val Glu Glu Leu Ala Gln Asp Ile Leu Ser Lys Leu Pro Arg Asp Phe
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Asp Leu Glu Glu Val Met Lys Leu Tyr Pro Val Val Tyr Glu Glu Ser
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Met Asn Thr Val Leu Arg Gln Glu Leu Ile Arg Phe Asn Arg Leu Thr
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Lys Val Val Arg Arg Ser Leu Ile Asn Leu Gly Arg Ala Ile Lys Gly
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Gln Val Leu Met Ser Ser Glu Leu Glu Glu Val Phe Asn Ser Met Leu
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Gly Phe Tyr Phe Thr Gln Ser Phe Leu Thr Gly Val Ser Gln Asn Tyr
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Ala Arg Lys Tyr Thr Ile Pro Ile Asp His Ile Gly Phe Glu Phe Glu
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Val Thr Pro Gln Glu Thr Val Met Glu Asn Asn Pro Glu Asp Gly Ala
                        855
                                            860
Tyr Ile Lys Gly Leu Phe Leu Glu Gly Ala Arg Trp Asp Arg Lys Thr
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                                        875
Met Gln Ile Gly Glu Ser Leu Pro Lys Ile Leu Tyr Asp Pro Leu Pro
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                                    890
Ile Ile Trp Leu Lys Pro Gly Glu Ser Ala Met Phe Leu His Gln Asp
                                905
 Ile Tyr Val Cys Pro Val Tyr Lys Thr Ser Ala Arg Arg Gly Thr Leu
                            920
 Ser Thr Thr Gly His Ser Thr Asn Tyr Val Leu Ser Ile Glu Leu Pro
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 Gly Ala Ala Ser Ala Val Ser Leu Ala Gly Ala Ser Leu Val Leu Ser
                                                 45
 Leu Leu Gln Arg Val Ala Ser Tyr Ala Arg Lys Trp Gln Gln Met Arg
                         55
 Pro Ile Pro Thr Val Ala Arg Ala Tyr Pro Leu Val Gly His Ala Leu
                                         75
                     70
 Leu Met Lys Pro Asp Gly Arg Glu Phe Phe Gln Gln Ile Ile Glu Tyr
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 Thr Glu Glu Tyr Arg His Met Pro Leu Leu Lys Leu Trp Val Gly Pro
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Val Pro Met Val Ala Leu Tyr Asn Ala Glu Asn Val Glu Val Ile Leu
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Thr Ser Ser Lys Gln Ile Asp Lys Ser Ser Met Tyr Lys Phe Leu Glu
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Pro Trp Leu Gly Leu Gly Leu Leu Thr Ser Thr Gly Asn Lys Trp Arg
                                         155
                    150
Ser Arg Arg Lys Met Leu Thr Pro Thr Phe His Phe Thr Ile Leu Glu
                                    170
Asp Phe Leu Asp Ile Met Asn Glu Gln Ala Asn Ile Leu Val Lys Lys
                                185
Leu Glu Lys His Ile Asn Gln Glu Ala Phe Asn Cys Phe Phe Tyr Ile
                            200
Thr Leu Cys Ala Leu Asp Ile Ile Cys Glu Thr Ala Met Gly Lys Asn
                        215
Ile Gly Ala Gln Ser Asn Asp Asp Ser Glu Tyr Val Arg Ala Val Tyr
                    230
                                         235
225
Arg Met Ser Glu Met Ile Phe Pro Arg Ile Lys Met Pro Trp Leu Trp
                                     250
                245
Leu Asp Leu Trp Tyr Leu Met Phe Lys Glu Gly Trp Glu His Lys Lys
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 ggattatteeg
 ctagccageg
 acatggggt

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 ggaacgaaga
 tecegaggag
 cetgcctacg
 gagacacggc

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 tecacgtggg
 accetetgeg
 gacagtggca
 gegagtett

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 teggtggata
 geggttgaag
 aactgcaca

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 tegatgggta
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 cettggecac

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Lys Glu Arg Pro Ser Arg Arg Ala Arg Gly Ser Pro Phe Val Arg Ser
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Gly Thr Ile Val Arg Ser Gln Thr Phe Ser Pro Gly Ala Arg Ser Gln
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                                        75
Tyr Val Cys Arg Leu Tyr Arg Ser Asp Ser Asp Ser Ser Thr Leu Pro
                                    90
Arg Lys Ser Pro Phe Val Arg Asn Thr Leu Glu Arg Arg Thr Leu Arg
                                105
                                                     110
Tyr Lys Gln Ser Cys Arg Ser Ser Leu Ala Glu Leu Met Ala Arg Thr
                                                125
                            120
Ser Leu Asp Leu Glu Leu Asp Leu Gln Ala Ser Arg Thr Arg Gln Arg
                        135
                                            140
Gln Leu Asn Glu Glu Leu Cys Ala Leu Arg Glu Leu Arg Gln Arg Leu
                    150
                                        155
Glu Asp Ala Gln Leu Arg Gly Gln Thr Asp Leu Pro Pro Trp Val Leu
Arg Asp Glu Arg Leu Arg Gly Leu Leu Arg Glu Ala Glu Arg Gln Thr
                                185
Arg Gln Thr Lys Leu Asp Tyr Arg His Glu Gln Ala Ala Glu Lys Met
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Leu Lys Lys Ala Ser Lys Glu Ile Tyr Gln Leu Arg Gly Gln Ser His
                       215
Lys Glu Pro Ile Gln Val Gln Thr Phe Arg Glu Lys Ile Ala Phe Phe
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Gln Gln Gln Arg Gln Arg Leu Ala Arg His Gly Val Arg Arg Ala Ala
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Pro Arg Arg Leu Val Val Leu Glu Asp Glu Val Glu Leu Asp Leu Gln
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                                           60
His Glu Asp Val Lys Glu Pro Gln Asp His Gly Val Ala Ala Leu Gly
                                                          80
                                       75
                    70
Arg Ala His Leu Gly Ala His Pro His Gly His Val Ala Gln His Gln
Gln Glu Ala His Val Ala His Gln His Asp Asp Ala His Ala Asp Leu
                               105
            100
Ala Arg Ala Leu Val Leu Leu His Gln Val Arg Val His Asp Gly His
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Ala Ala His Asp His Gln Arg Gly Gln Ala His Val Ala Pro Val Arg
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                                 25
Glu Ser Gly Leu Gly Lys Ser Thr Leu Ile Asn Ser Leu Phe Leu Thr
                            40
Asn Leu Tyr Glu Asp Arg Gln Val Pro Glu Ala Ser Ala Arg Leu Thr
                                            60
Gln Thr Leu Ala Ile Glu Arg Arg Gly Val Glu Ile Glu Glu Gly Gly
Val Lys Val Lys Leu Thr Leu Val Asp Thr Pro Gly Phe Gly Asp Ser
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90
Val Asp Cys Ser Asp Cys Trp Leu Pro Val Val Lys Phe Ile Glu Glu
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            100
Gln Phe Glu Gln Tyr Leu Arg Asp Glu Ser Gly Leu Asn Arg Lys Asn
                            120
        115
Ile Gln Asp Ser Arg Val His Cys Cys Leu Tyr Phe Ile Ser Pro Phe
                                            140
                        135
Gly Arg Ala Pro Ala Pro Arg Cys Gly Phe Leu Arg Ala Ile His Glu
                                        155
                    150
Lys Val Asn Ile Ile Pro Val Ile Gly Lys Ala Asp Ala Leu Met Pro
                165
                                    170
Gln Glu Thr Gln Ala Leu Lys Gln Lys Ile Arg Asp Gln Leu Lys Glu
                                185
            180
Glu Glu Ile His Ile Tyr Gln Phe Pro Glu Cys Asp Ser Asp Glu Asp
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                            200
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Glu Asp Phe Lys Arg Gln Asp Ala Glu Met Lys Glu Ser Ile Pro Phe
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Ala Val Val Gly Ser Cys Glu Val Val
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ggcgccggtg gtcgttgtga cccaacctgg agtcggtccc ggtccggccc cccagaactc
180
caactggcag acaggcatgt gtgactgttt cagcgactgc ggagtctgtc tctgtggcac
attttgtttc ccgtgccttg ggtgtcaagt tgcagctgat atgaatgaat gctgtctgtg
tggaacaagc gtcgcaatga ggactctcta caggacccga tatggcatcc ctggatctat
360
ttgtgatgac tatatggcaa ctctttgctg tcctcattgt actctttgcc aaatcaagag
agatatcaac agaaggagag ccatgcgtac tttctaaaaa ctgatggtga aaagctctta
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 720
 tataaaccaa atgaaatatt ttactgataa gattcttcat gcttctttgc tctccttaaa
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840

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864
<210> 4660
<211> 192
<212> PRT
<213> Homo sapiens
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Met Pro Ser Val Val Leu Lys His Ile His His Ile Ser Val Ala Lys
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Asp Gly Glu Glu Leu Lys Leu Lys Arg Cys Leu Leu Asn Phe Val Ala
                                25
            20
Ser Val Arg Ala Phe His His Gln Phe Leu Glu Ser Thr His Gly Ser
        35
                            40
                                                 45
Pro Ser Val Asp Ile Ser Leu Asp Leu Ala Lys Ser Thr Met Arg Thr
                        55
Ala Lys Ser Cys His Ile Val Ile Thr Asn Arg Ser Arg Asp Ala Ile
                    70
Ser Gly Pro Val Glu Ser Pro His Cys Asp Ala Cys Ser Thr Gln Thr
                                    90
Ala Phe Ile His Ile Ser Cys Asn Leu Thr Pro Lys Ala Arg Glu Thr
                                105
Lys Cys Ala Thr Glu Thr Asp Ser Ala Val Ala Glu Thr Val Thr His
                            120
Ala Cys Leu Pro Val Gly Val Leu Gly Gly Arg Thr Gly Thr Asp Ser
                        135
                                             140
Arg Leu Gly His Asn Asp His Arg Arg Leu Ser Leu His Phe Gln Cys
                    150
                                         155
145
Arg Ala Phe His Val Val Phe Ile Cys Gly Glu Ile Leu Ser Gln Ala
                                     170
Thr Arg His Phe Leu Leu Gly Thr Leu Phe Thr Asn Phe His Cys Phe
                                185
<210> 4661
<211> 153
<212> DNA
<213> Homo sapiens
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tttgaggacc ctcaccatgg ccatgggcag ttc
153
<210> 4662
<211> 51
<212> PRT
<213> Homo sapiens
<400> 4662
Arg Ile Cys Met Pro Leu Thr Val Asp Glu Tyr Lys Ile Gly Gln Leu
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Tyr Met Ile Ser Lys His Ser His Glu Gln Ser Asp Arg Gly Glu Gly
            20
                                25
Val Glu Val Val Gln Asn Glu Pro Phe Glu Asp Pro His His Gly His
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Gly Gln Phe
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<211> 1550
<212> DNA
<213> Homo sapiens
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cagacggatg acccaggece cetegatgge cetgacetee aggecageca etcagagete
caggtgccca cccctggcag agccggccta ctgaacacct ctggtaccaa aggcttagaa
tgttctcctt caactcccac catgaattct tacttttata agttcatgat caaccttctc
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ctgtgcctcc tgctgaatgc ggagaacatc ttccactcaa tggcagacat cctgctgcgg
gaggaggacc tcaagttcgc ctcgaccatg gtccacgccc tcaacaccat cctgctgacc
tccacagage tettecaget aaggaaccag etgaaggace tgaagaccet ggagagccag
aacctgttct gctgcctgta ccgctcctgg tgccacaacc cagtcaccac ggtgtccctc
tgcttcctca cccagaacta ccggcacgcc tatgacctca tccagaagtt tggggacctg
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720
atottcacat atotgogoot goagotgotg gaogtgaaga acaacccota cotgatcaag
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cggctccagt gcgtgcccaa ccctgagctg ctgcagaccg aagacagtct aaaggcagcc
cccaagtccc agaaagctga ctcccctagc atcgactacg cagagctgct gcagcacttt
qaqaaggtcc agaacaagca cctggaagtg cggcaccagc ggagcgggcg tggggaccac
ctggaccgga gggttgtcct ctgacaggcc tggcacggag gagggcccac cgagtggtcc
catgaaacac taagggtegt cacgccctcc cgaggagctc aaggacctgc ctgtcaggac
cagggctggg cctgccaacc cagggcagtg ttggggccgg aggctgctgt gtctgcccaa
1200
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getectetea gagtecagte eccaggeete eagegetgte agetgeacce tggcattete
acagagetgg etgeceacce agtgggggge tatageetca gagaceacte atcetetgga
atcaacctct ttctaatacc ctcttggaaa aagagcttgc ccctcctcca gcacactaga
gctctggcct tgtgtgtata tgtatacata cgtgaacaca tgcctgtgtg tgtgtgtgt
tgtgtacttg tatgcacgta ggcaccagca caaagatctg aatgatgcac cccacccca
<210> 4664
<211> 347
<212> PRT
<213> Homo sapiens
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Glu Ile Ala Ser Ser Pro Ala Gly Gln Thr Asp Asp Pro Gly Pro Leu
Asp Gly Pro Asp Leu Gln Ala Ser His Ser Glu Leu Gln Val Pro Thr
                       55
Pro Gly Arg Ala Gly Leu Leu Asn Thr Ser Gly Thr Lys Gly Leu Glu
                                       75
                                                          80
                   70
Cys Ser Pro Ser Thr Pro Thr Met Asn Ser Tyr Phe Tyr Lys Phe Met
Ile Asn Leu Leu Lys Arg Phe Ser Ser Glu Arg Lys Leu Leu Glu Val
                               105
Arg Gly Pro Phe Ile Ile Arg Gln Leu Cys Leu Leu Asn Ala Glu
                           120
        115
Asn Ile Phe His Ser Met Ala Asp Ile Leu Leu Arg Glu Glu Asp Leu
                       135
                                           140
Lys Phe Ala Ser Thr Met Val His Ala Leu Asn Thr Ile Leu Leu Thr
                   150
                                       155
Ser Thr Glu Leu Phe Gln Leu Arg Asn Gln Leu Lys Asp Leu Lys Thr
               165
                                   170
Leu Glu Ser Gln Asn Leu Phe Cys Cys Leu Tyr Arg Ser Trp Cys His
                               185
Asn Pro Val Thr Thr Val Ser Leu Cys Phe Leu Thr Gln Asn Tyr Arg
                           200
        195
His Ala Tyr Asp Leu Ile Gln Lys Phe Gly Asp Leu Glu Val Thr Val
                                           220
                       215
Asp Phe Leu Ala Glu Val Asp Lys Leu Val Gln Leu Ile Glu Cys Pro
                   230
                                       235
Ile Phe Thr Tyr Leu Arg Leu Gln Leu Leu Asp Val Lys Asn Asn Pro
               245
                                   250
Tyr Leu Ile Lys Ala Leu Tyr Gly Leu Leu Met Leu Leu Pro Gln Ser
                               265
Ser Ala Phe Gln Leu Leu Ser His Arg Leu Gln Cys Val Pro Asn Pro
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285
                           280
       275
Glu Leu Leu Gln Thr Glu Asp Ser Leu Lys Ala Ala Pro Lys Ser Gln
                       295
Lys Ala Asp Ser Pro Ser Ile Asp Tyr Ala Glu Leu Leu Gln His Phe
                                      315
                   310
Glu Lys Val Gln Asn Lys His Leu Glu Val Arg His Gln Arq Ser Gly
               325
                                  330
Arg Gly Asp His Leu Asp Arg Arg Val Val Leu
                               345
           340
<210> 4665
<211> 1043
<212> DNA
<213> Homo sapiens
<400> 4665
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agagtettea tgtggacagt eteagggaca ecatgtagag aattttggte tegatteaga
aaagagaaag agccagtggt tgttgagaca gtagaagaga aaaaggaacc tatcctagtg
tqtccacctt tacgaagccg agcatacaca ccacctgaag atctccagag tcgtttggaa
tettacgtta aagaagtttt tggttcatet etteetagta attggcaaga cateteeetg
gaagatagtc qtctaaagtt caatcttctg gctcatttag ctgatgactt gggtcatgta
360
gtccctaact ccagactcca ccagatgtgc agggttagag atgttcttga tttctataat
gtccctattc aagatagatc taaatttgat gaactcaqtg ccagtaatct gccccccaat
ttqaaaatca cttggaqtta ctaagcaatt cggaagagaa acacattgaa atcactgtct
ttccctgagc aagggggctg ctcattagat cttttgatac tttaccatgt gaaatactac
cagaactgtt ctctaaaccc actttttctg tagaggaatg tatcatcttt tttttctca
tattacaaat ggacaaataa cggactttct attttcatat ttgctgaaac cattttttaa
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ttcatcctgt taggattcat atctaagata gagttatgca ttgcacatac acaaataaac
ttttattaga tagataccta taaaagaaac ataaaagtat gttgtgtatt actgacaqtt
aaaaaaaaa aaaaaaaaa aaa
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<210> 4666

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<211> 167
<212> PRT
<213> Homo sapiens
<400> 4666
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Gly Ile Thr Arg Arg Val Phe Met Trp Thr Val Ser Gly Thr Pro Cys
                                25
Arg Glu Phe Trp Ser Arg Phe Arg Lys Glu Lys Glu Pro Val Val Val
                            40
Glu Thr Val Glu Glu Lys Lys Glu Pro Ile Leu Val Cys Pro Pro Leu
                        55
Arg Ser Arg Ala Tyr Thr Pro Pro Glu Asp Leu Gln Ser Arg Leu Glu
                                        75
                    70
Ser Tyr Val Lys Glu Val Phe Gly Ser Ser Leu Pro Ser Asn Trp Gln
                                    90
Asp Ile Ser Leu Glu Asp Ser Arg Leu Lys Phe Asn Leu Leu Ala His
                                105
            100
Leu Ala Asp Asp Leu Gly His Val Val Pro Asn Ser Arg Leu His Gln
                            120
Met Cys Arg Val Arg Asp Val Leu Asp Phe Tyr Asn Val Pro Ile Gln
                        135
Asp Arg Ser Lys Phe Asp Glu Leu Ser Ala Ser Asn Leu Pro Pro Asn
                                        155
                                                             160
145
Leu Lys Ile Thr Trp Ser Tyr
                165
<210> 4667
<211> 1031
<212> DNA
<213> Homo sapiens
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cctctgctgg aggggaaagc ccgctcctgt tttgctatga ccgagcccca ggttgcctct
tcaqatgcca ccaacattga ggcttccatc agagaggagg acagcttcta tgtcataaac
ggtcacaaat ggtggatcac aggcatcctg gatcctcgtt gccaactctg tgtgtttatg
ggaaaaacag acccacatgc accaagacac cggcagcagt ctgtgctctt ggttcccatg
gataccccag ggataaaaat catccggcct ctgacggtgt atggactgga agatgcacca
ggtggccatg gtgaagtccg atttgagcac gtgcgtgtgc ccaaagagaa catggtcctg
ggccctggcc gaggctttga gatcgcccag ggcagactgg gccccggcag gatccatcac
tgcatgaggc tgatcgggtt ctcagagagg gccctggcac tcatgaaggc ccgcgtgagt
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gettteecce geacceagea etgacteaga accaecacet tetgetttge tgteggactt
caatteetae etgttttetg agtgeagtee tageaggtga ageaaggtga tgteettgee
aagaagttgc attectgtct getttgcate tgctactttg etgcagtttg gattcagage
agaatggacc ccactctgtc gaggtgacct gaagggaaac gccaggctct gtagcagcag
agggcaaggt tccaaggtgt aaaggtcatg ctgctagcac attattaaaa atcagtctgg
gtgcaatggc tcacagctat aatcccagta ctttgggagg tctaggtagg agggttgctt
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aaaaaaaaa a
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<210> 4668
<211> 207
<212> PRT
<213> Homo sapiens
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Xaa Ala Met Gly Thr Ser Leu Tyr Ala Pro Glu Val Cys Asn Cys Ser
                                     10
Ala Pro Asp Thr Gly Asn Met Glu Leu Leu Val Arg Tyr Gly Thr Glu
                                 25
 Ala Gln Lys Ala Arg Trp Leu Ile Pro Leu Leu Glu Gly Lys Ala Arg
                             40
 Ser Cys Phe Ala Met Thr Glu Pro Gln Val Ala Ser Ser Asp Ala Thr
 Asn Ile Glu Ala Ser Ile Arg Glu Glu Asp Ser Phe Tyr Val Ile Asn
 Gly His Lys Trp Trp Ile Thr Gly Ile Leu Asp Pro Arg Cys Gln Leu
                                     90
 Cys Val Phe Met Gly Lys Thr Asp Pro His Ala Pro Arg His Arg Gln
             100
                                 105
 Gln Ser Val Leu Leu Val Pro Met Asp Thr Pro Gly Ile Lys Ile Ile
                                                 125
                             120
 Arg Pro Leu Thr Val Tyr Gly Leu Glu Asp Ala Pro Gly Gly His Gly
                         135
                                             140
 Glu Val Arg Phe Glu His Val Arg Val Pro Lys Glu Asn Met Val Leu
                                         155
                     150
 Gly Pro Gly Arg Gly Phe Glu Ile Ala Gln Gly Arg Leu Gly Pro Gly
                                      170
                 165
 Arg Ile His His Cys Met Arg Leu Ile Gly Phe Ser Glu Arg Ala Leu
                                  185
 Ala Leu Met Lys Ala Arg Val Ser Ala Phe Pro Arg Thr Gln His
         195
                             200
 <210> 4669
 <211> 683
 <212> DNA
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<213> Homo sapiens

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cagaaagtat ttcaacacaa tgaacttaag aaagagactt gtgttcaggc aqgttttcag
120
qacatqaaca taaaaaaaca gattcaggaa cagcaccagg ctgccattat tattcagaag
cattgtaaag cctttaaaat aaggaagcat tatctccaca ttagagcaac agtagtttct
attcaaagaa gatacagaaa actaactgca gtgcgtaccc aagcagttat ttgtatacag
300
tottattaca gaggotttaa agtacgaaag gatattcaaa atatgcaccg ggotgccaca
ctaattcagt cattctatcg aatgcacagg gccaaagttg attattaaac aaagaaaact
gcaattgtgg ttatacagaa ttattatagg ttgtatgtta gagtaaaaac agaaagaaaa
aactttttag cagttcagaa atctgtccga actattcagg ctgcttttag aggcatgaaa
gttagacaaa aattgaaaaa atgtatcaga ggaaaagatg gcagccattg ttaaccaatc
tgcactctgc tgttacagaa gtaaaactca gtatgaagct gttcaaagtg aaggtgttat
gattcaagag tggtataaag ctt
683
<210> 4670
<211> 135
 <212> PRT
 <213> Homo sapiens
 <400> 4670
 Xaa Ser Phe Ser Gly Leu Arg Gly Ile Ile Gln Glu Lys Tyr Arg Ala
 Asn Lys Lys Lys Gln Lys Val Phe Gln His Asn Glu Leu Lys Lys Glu
 Thr Cys Val Gln Ala Gly Phe Gln Asp Met Asn Ile Lys Lys Gln Ile
 Gln Glu Gln His Gln Ala Ala Ile Ile Ile Gln Lys His Cys Lys Ala
 Phe Lys Ile Arg Lys His Tyr Leu His Ile Arg Ala Thr Val Val Ser
 Ile Gln Arg Arg Tyr Arg Lys Leu Thr Ala Val Arg Thr Gln Ala Val
 Ile Cys Ile Gln Ser Tyr Tyr Arg Gly Phe Lys Val Arg Lys Asp Ile
                                 105
 Gln Asn Met His Arg Ala Ala Thr Leu Ile Gln Ser Phe Tyr Arg Met
                             120
         115
 His Arg Ala Lys Val Asp Tyr
                         135
     130
 <210> 4671
 <211> 657
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<212> DNA
<213> Homo sapiens
<400> 4671
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gcaccagege cateegette gaggttgage eteetgeage agtggaatea ggggeeteet
ggggetegge aggggetace eggeteeget teegeecagt aatggagaet geagecaegt
taggccaggc tgctgcagtg gtttcagcat ctatccgcag ggatccacgg ggaagctggt
gtgcgccgga taaagatggc aaccgccgat gagattgtga aactcatgct cgaccacatg
300
acaaacacca ccaacgcgtc ccatgtgcct gtgcagcccg gctcctcagt tgtgatqatq
gtcaacaacc tgggtggcct gtcattcctg gaactgggca tcatagccga cgctaccgtc
cgctccctgg agggccgcgg ggtgaagatt gcccgtgccc tggtgggcac cttcatgtca
quantification of the control of the
ctgatagatg ctgaaaccac tgcagcagcc tggcctcgaa geggatggcg ctggtgctgg
aacgggtgtg cagcactctc ctgggcctgg aggaacacct gaatgccctg gaccggt
<210> 4672
<211> 152
<212> PRT
<213> Homo sapiens
<400> 4672
Ala Arg Leu Leu Gln Trp Phe Gln His Leu Ser Ala Gly Ile His Gly
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Glu Ala Gly Val Arg Arg Ile Lys Met Ala Thr Ala Asp Glu Ile Val
                                                                             25
 Lys Leu Met Leu Asp His Met Thr Asn Thr Thr Asn Ala Ser His Val
                                                                   40
 Pro Val Gln Pro Gly Ser Ser Val Val Met Met Val Asn Asn Leu Gly
                                                          55
 Gly Leu Ser Phe Leu Glu Leu Gly Ile Ile Ala Asp Ala Thr Val Arg
                                                                                                75
                                                70
 Ser Leu Glu Gly Arg Gly Val Lys Ile Ala Arg Ala Leu Val Gly Thr
                                                                                      90
 Phe Met Ser Ala Leu Glu Met Pro Gly Ile Ser Leu Thr Leu Leu Leu
                                                                                                                            110
                                                                             105
 Val Asp Glu Pro Leu Leu Lys Leu Ile Asp Ala Glu Thr Thr Ala Ala
                                                                   120
                                                                                                                   125
 Ala Trp Pro Arg Ser Gly Trp Arg Trp Cys Trp Asn Gly Cys Ala Ala
                                                          135
                                                                                                         140
 Leu Ser Trp Ala Trp Arg Asn Thr
                                                150
 145
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<211> 1335 <212> DNA <213> Homo sapiens <400> 4673 cegeggette tggetgegeg geetgegege geeteeeggg eggatteeag eecegagegg gacagegeag eggggagega egagatttet etetgateaa aeggaeagtt eaggaeteag aatctaagga tgaatgttca ccgtggcagt gacagtgaca ggttattgcg gcaggaggcc agetgettag tggatgatae tttagetgta geccaagaaa aagaageaaa cageetgget teatetggte eteataatet taettateet etaggteeca ggaatgaaga eeteteaett gactatgcct ctcagccagc aaatcttcag ttccctcaca taatgcccct tgctgaagac atcaaaggtt cttgcttcca aagtgggaat aaacggaacc atgaaccttt tattgctcca gaaagatttg gaaacagtag tgtgggcttt ggcagtaatt cccattccca agcaccagag aaagtgacgc ttcttgtaga tggcacacgt tttgttgtga atccacagat tttcactgct catccggata ccatgctggg aaggatgttt ggaccaggaa gagagtacaa cttcactcgg cccaatgaga agggagagta tgagattgct gaaggcatca gtgcaactgt atttcgcaca gtgctggatt attacaaaac cggtatcatc aattgtcctg atggcatctc tatcccagat cttagagata cttgtgatta tctctgcatt aattttgact tcaacactat ccgatgtcaa gatctgagtg ctttactcca tgaactgtct aatgacggtg ctcataagca gtttgatcac tacctcgaag agctcatctt gcccatcatg gtgggctgtg ccaagaaagg agaacgagag tqccacattg ttgtgctgac ggatgaggat tctgtggact gggatgaaga ccaccctcca ccaatggggg aggaatattc ccaaattctt tatagctcca agctctacag attcttcaaa tatattgaga atagggatgt tgcaaaaaca gtgttaaagg aacggggcct aaaaaacatt cgcattggaa ttgaaggtta ccctacctgt aaagaaaaa ttaagagaag gcctggcggc egttetgaag teatetataa ttatgtacaa egeceettea teeagatgte atgggaaaag gaagaaggga agagtegeca tgtggattte cagtgtgtte gaagcaaate cetcacgaat ctggtagctg ctggagatga tgtcttggag gaccaggaga tattaatgca tcacccaccc caagtggatg aactt 1335

<210> 4674

<210> 4673

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<211> 402
<212> PRT
<213> Homo sapiens
<400> 4674
Met Asn Val His Arg Gly Ser Asp Ser Asp Arg Leu Leu Arg Gln Glu
                                    10
Ala Ser Cys Leu Val Asp Asp Thr Leu Ala Val Ala Gln Glu Lys Glu
                                2.5
            20
Ala Asn Ser Leu Ala Ser Ser Gly Pro His Asn Leu Thr Tyr Pro Leu
                            40
Gly Pro Arg Asn Glu Asp Leu Ser Leu Asp Tyr Ala Ser Gln Pro Ala
                        55
Asn Leu Gln Phe Pro His Ile Met Pro Leu Ala Glu Asp Ile Lys Gly
                    70
                                        75
Ser Cys Phe Gln Ser Gly Asn Lys Arg Asn His Glu Pro Phe Ile Ala
                                    90
Pro Glu Arg Phe Gly Asn Ser Ser Val Gly Phe Gly Ser Asn Ser His
                                105
            100
Ser Gln Ala Pro Glu Lys Val Thr Leu Leu Val Asp Gly Thr Arg Phe
                            120
                                                125
Val Val Asn Pro Gln Ile Phe Thr Ala His Pro Asp Thr Met Leu Gly
                                            140
                        135
Arg Met Phe Gly Pro Gly Arg Glu Tyr Asn Phe Thr Arg Pro Asn Glu
                    150
                                        155
Lys Gly Glu Tyr Glu Ile Ala Glu Gly Ile Ser Ala Thr Val Phe Arg
                                    170
                165
Thr Val Leu Asp Tyr Tyr Lys Thr Gly Ile Ile Asn Cys Pro Asp Gly
                                185
            180
 Ile Ser Ile Pro Asp Leu Arg Asp Thr Cys Asp Tyr Leu Cys Ile Asn
                            200
Phe Asp Phe Asn Thr Ile Arg Cys Gln Asp Leu Ser Ala Leu Leu His
                        215
Glu Leu Ser Asn Asp Gly Ala His Lys Gln Phe Asp His Tyr Leu Glu
                                         235
                    230
Glu Leu Ile Leu Pro Ile Met Val Gly Cys Ala Lys Lys Gly Glu Arg
                245
                                    250
Glu Cys His Ile Val Val Leu Thr Asp Glu Asp Ser Val Asp Trp Asp
                                 265
Glu Asp His Pro Pro Pro Met Gly Glu Glu Tyr Ser Gln Ile Leu Tyr
                                                285
                            280
 Ser Ser Lys Leu Tyr Arg Phe Phe Lys Tyr Ile Glu Asn Arg Asp Val
                                            300
                        295
 Ala Lys Thr Val Leu Lys Glu Arg Gly Leu Lys Asn Ile Arg Ile Gly
                                        315
                     310
 Ile Glu Gly Tyr Pro Thr Cys Lys Glu Lys Ile Lys Arg Arg Pro Gly
                                     330
 Gly Arg Ser Glu Val Ile Tyr Asn Tyr Val Gln Arg Pro Phe Ile Gln
                                345
 Met Ser Trp Glu Lys Glu Glu Gly Lys Ser Arg His Val Asp Phe Gln
                            360
 Cys Val Arg Ser Lys Ser Leu Thr Asn Leu Val Ala Ala Gly Asp Asp
                         375
 Val Leu Glu Asp Gln Glu Ile Leu Met His His Pro Pro Gln Val Asp
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385 390 395 400 Glu Leu

<210> 4675

<211> 2868 <212> DNA

<213> Homo sapiens

<400> 4675

1320

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Asn Ser Phe Cys Ser Asp Asp Thr Gly Cys Pro Ser Ser Gln Ser Val
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Ser Pro Val Lys Thr Pro Ser Asp Ala Gly Asn Ser Pro Ile Gly Phe
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Cys Pro Gly Ser Asp Glu Gly Phe Thr Arg Lys Lys Cys Thr Ile Gly
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Met Val Gly Glu Gly Ser Ile Gln Ser Ser Arg Tyr Lys Lys Glu Ser
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Lys Ser Gly Leu Val Lys Pro Gly Ser Glu Ala Asp Phe Ser Ser
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Ser Ser Thr Gly Ser Ile Ser Ala Pro Glu Val His Met Ser Thr Ala
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Gly Ser Lys Arg Ser Ser Ser Ser Arg Asn Arg Gly Pro His Gly Arg
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Ser Asn Gly Ala Ser Ser His Lys Pro Gly Ser Ser Ser Ser Pro
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Arg Glu Lys Asp Leu Leu Ser Met Leu Cys Arg Asn Gln Leu Ser Pro
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Val Asn Ile His Pro Ser Tyr Ala Pro Ser Ser Pro Ser Ser Ser Asn
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Ser Gly Ser Tyr Lys Gly Ser Asp Cys Ser Pro Ile Met Arg Arg Ser
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Gly Arg Tyr Met Ser Cys Gly Glu Asn His Gly Val Arg Pro Pro Asn
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Pro Glu Gln Tyr Leu Thr Pro Leu Gln Gln Lys Glu Val Thr Val Arg
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His Leu Lys Thr Lys Leu Lys Glu Ser Glu Arg Arg Leu His Glu Arg
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Glu Ser Glu Ile Val Glu Leu Lys Ser Gln Leu Ala Arg Met Arg Glu
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Asp Trp Ile Glu Glu Glu Cys His Arg Val Glu Ala Gln Leu Ala Leu
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Lys Glu Ala Arg Lys Glu Ile Lys Gln Leu Lys Gln Val Ile Glu Thr
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Val Asp Ile Asn Ile Gln Asn Lys Lys Leu Glu Ser Leu Leu Gln Ser
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Met Glu Met Ala His Ser Gly Ser Leu Arg Asp Glu Leu Cys Leu Asp
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Phe Pro Cys Asp Ser Pro Glu Lys Ser Leu Thr Leu Asn Pro Pro Leu
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Asp Thr Met Ala Asp Gly Leu Ser Leu Glu Glu Gln Val Thr Gly Glu
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Asp Leu Phe Asp Glu Ile Val Thr Ala Thr Thr Thr Glu Ser Gly Asp
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Leu Glu Leu Val His Ser Thr Pro Gly Ala Asn Val Leu Glu Leu Leu
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Pro Ile Val Met Gly Gln Glu Glu Gly Ser Val Val Val Glu Arg Ala
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Gln Ser Val Leu Gln Lys Leu Gln Asp Pro Cys Pro Ser Ser Leu Ala
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Ser Pro Asp Glu Ser Glu Pro Asp Ser Met Glu Ser Phe Pro Glu Ser
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Leu Ser Ala Leu Val Val Asp Leu Thr Pro Arg Asn Pro Asn Ser Ala
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Ile Leu Leu Ser Pro Val Glu Thr Pro Tyr Xaa Gln Cys Gly Cys Arg
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Ser Ser Cys Lys Pro Pro His Glu Arg Ala Gly Xaa Phe Ala Ala Cys
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Val Glu Glu Arg Leu Asp Gly Val Ile Pro Leu Ala Arg Gly Gly Val
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Val Arg Gln Tyr Trp Ser Ser Ser Phe Leu Val Asp Leu Leu Ala Val
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Ala Ala Pro Val Val Pro Thr Val Leu Trp Ala Phe Ser Thr Gln Arg
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Gly Gly Thr Asp Pro Val Tyr Asn Ile Gly Ala Leu Leu Arg Gly Cys
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Cys Val Val Ala Leu His Ser Leu Arg Arg Thr Ala Phe Arg Ile Lys
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Thr
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Arg Thr Val Phe Ile Trp Phe Val Gly Gln Leu Leu Gly Gly Glu Leu
Lys Gly Tyr Ser Lys Thr Asn Thr Thr Ser Ser Arg Pro Ala Ser Ser
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Arg Gly Ser Leu Ser Ser Ser Ser Ser Ser Ser Ser Leu Thr Lys
                    70
Asp Ala Leu Pro Ser Ser Leu Lys Ser Asp Ser Thr Thr Ile Thr Ser
                85
                                    90
Gly Leu Val Phe Pro Phe Arg Ser Leu Cys Val Asn Pro Ala Lys Ser
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Ser Val Ser Glu Ser Val Ser Ser Ile Lys Ile Leu Leu Ser Ser Ser
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Ser Pro Cys Ser Leu Thr Phe Ser Arg Ala Ile Lys Ala Thr Ser Ser
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Ile Ala Gly Pro Gln Thr Phe Gln Gly Lys His Cys Phe Thr Ser Cys
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Arg Gln Leu Ile Ser Gln Lys Pro Leu Gln Lys Pro Val Leu Pro Gly
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Phe Leu Phe His Gln Thr Thr Arg Gln Lys Asn Leu Ser Phe Leu Pro
Pro Phe Ser Phe Phe Pro Ser Cys Thr His Leu Glu Asn Phe Thr Phe
Leu Glu Ser Pro Gln Asn Asn Thr Lys Val Ile Val Gly Ala Thr Gly
Phe Met Leu Tyr Cys Gly Ala Arg Gly Lys Thr Cys Leu Tyr Ala Gly
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Asn Thr His Asn His Ser Phe Arg Phe Val Cys Leu Met Val Ile Cys
His Lys Arg Asp Leu Gln Lys Gln Gly Ala Leu Val Asn Val Gln Tyr
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Leu Asp Phe Cys Val Leu Arg Thr Gln Lys Gly Ala Thr Leu Leu Phe
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Gly Pro Val Ser Gly His Leu Val Ile
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Gln Thr His Gly Thr Ala Arg Ile Gly Thr His Asn Gly Thr Phe His
Cys Asp Glu Ala Leu Ala Cys Ala Leu Leu Arg Leu Leu Pro Glu Tyr
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Arg Asp Ala Glu Ile Val Arg Thr Arg Asp Pro Glu Lys Leu Ala Ser
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Cys Asp Ile Val Val Asp Val Gly Glu Tyr Asp Pro Arg Arg His
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Arg Tyr Asp His His Gln Arg Ser Phe Thr Glu Thr Met Ser Ser Leu
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Ser Pro Gly Lys Pro Trp Gln Thr Lys Leu Ser Ser Ala Gly Leu Ile
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Tyr Leu His Phe Gly His Lys Leu Leu Ala Gln Leu Leu Gly Thr Ser
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Glu Glu Asp Ser Met Val Gly Thr Leu Tyr Asp Lys Met Tyr Glu Asn
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Phe Val Glu Glu Val Asp Ala Val Asp Asn Gly Ile Ser Gln Trp Ala
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Glu Gly Glu Pro Arg Tyr Ala Leu Thr Thr Thr Leu Ser Ala Arg Val
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Ala Arg Leu Asn Pro Thr Trp Asn His Pro Asp Gln Asp Thr Glu Ala
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Gly Phe Lys Arg Ala Met Asp Leu Val Gln Glu Glu Phe Leu Gln Arg
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Leu Asp Phe Tyr Gln His Ser Trp Leu Pro Ala Arg Ala Leu Val Glu
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                                    250
Glu Ala Leu Ala Gln Arg Phe Gln Val Asp Pro Ser Gly Glu Ile Val
                                265
Glu Leu Ala Lys Gly Ala Cys Pro Trp Lys Glu His Leu Tyr His Leu
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Asp Gln Ala Gly Gln Trp Arg Ile Gln Cys Val Pro Lys Glu Pro His
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Ser Met Ala Arg Ala Thr Leu Ala Gln Arg Ser Tyr Leu Pro Gln Ile
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geogeagetg atgeoogagg acgogotgga cacoggtotg cagoogotto caacototoo
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tacctgcagt caaaggtggt ccgcgcaaag gagcgcctgg atgaggaact caaaatccag
gcccaggagg acagagaaaa agggcagatg ccccatacgt gactgctcgg ctccccccgc
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Gly Leu Ser Asp His Pro His Val His Thr Ala Ser Arg Ala Ala Ala
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Asp Ala Arg Gly Arg Ala Gly His Arg Ser Ala Ala Ala Ser Asn Leu
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                                 25
 Ser Gly Leu Ser Leu Gln Glu Ala Gln Gln Ile Leu Asn Val Ser Lys
Leu Ser Pro Glu Glu Val Gln Lys Asn Tyr Glu His Leu Phe Lys Val
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50
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Asn Asp Lys Ser Val Gly Gly Ser Phe Tyr Leu Gln Ser Lys Val Val
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Arg Ala Lys Glu Arg Leu Asp Glu Glu Leu Lys Ile Gln Ala Gln Glu
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Asp Arg Glu Lys Gly Gln Met Pro His Thr
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<212> DNA
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eggegetete geacecectg tgggtggcat tgatgagege cetaatectg ggtetgettt
tegtggeggt ctacagettg teccatggeg aggtetecta tgacecacte tatgetgget
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atggggggg
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<211> 90
<212> PRT
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Ala Leu Pro Val Ser Tyr Ala Leu Asn His Val Ser Ala Leu Ser His
Pro Leu Trp Val Ala Leu Met Ser Ala Leu Ile Leu Gly Leu Leu Phe
                            40
Val Ala Val Tyr Ser Leu Ser His Gly Glu Val Ser Tyr Asp Pro Leu
Tyr Ala Gly Phe Ala Val Phe Ala Phe Thr Ser Gly Gly Asp Leu Ile
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                    70
                                        75
Ile Ala Leu Gln Glu Asp Ser Tyr Gly Gly
<210> 4689
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ctgctggaca gctcagccag cgtctctcac tacgagttet cccgggttcg ggagtttgtg
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ctgtcagccg ctgcctcagc ccctgccgag aagcacctgc actttgtgga cgtggatgac
ctgcacatca ttgtccaaga gctgagggc tccattctcg acgcgatgcg gccacagcag
660
ctccatgcca cggagatcac gtccagcggc ttccgcctgg cctggccacc cctgctgacc
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cgccagcagc tgccagggaa cgccacggac tggatctggg ccggcctcga cccggacacg
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 <212> PRT
 <213> Homo sapiens
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 Ser Ala Pro Glu Asp Leu Met Phe Leu Leu Asp Ser Ser Ala Ser Val
 Ser His Tyr Glu Phe Ser Arg Val Arg Glu Phe Val Gly Gln Leu Val
 Ala Pro Leu Pro Leu Ala Pro Xaa Ala Leu Arg Ala Ser Leu Val His
                     70
                                          75
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 Val Gly Ser Arg Pro Tyr Thr Glu Phe Pro Phe Gly Gln His Ser Ser
                                      90
 Gly Glu Ala Ala Gln Asp Ala Val Arg Ala Ser Ala Gln Arg Met Gly
             100
                                 105
 Asp Thr His Thr Gly Leu Ala Leu Val Tyr Ala Lys Glu Gln Leu Phe
                             120
                                                  125
 Ala Glu Ala Ser Gly Ala Arg Pro Gly Val Pro Lys Val Leu Val Trp
                                              140
                         135
 Val Thr Asp Gly Gly Ser Ser Asp Pro Val Gly Pro Pro Met Gln Glu
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155
                    150
145
Leu Lys Asp Leu Gly Val Thr Val Phe Ile Val Ser Thr Gly Arg Gly
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                165
Asn Phe Leu Glu Leu Ser Ala Ala Ser Ala Pro Ala Glu Lys His
            180
                                185
Leu His Phe Val Asp Val Asp Asp Leu His Ile Ile Val Gln Glu Leu
                            200
                                                205
Arg Gly Ser Ile Leu Asp Ala Met Arg Pro Gln Gln Leu His Ala Thr
                        215
Glu Ile Thr Ser Ser Gly Phe Arg Leu Ala Trp Pro Pro Leu Leu Thr
                                        235
                    230
225
Ala Asp Ser Gly Tyr Tyr Val Leu Glu Leu Val Pro Ser Ala Gln Pro
                                     250
Gly Ala Ala Arg Arg Gln Gln Leu Pro Gly Asn Ala Thr Asp Trp Ile
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                                265
Trp Ala Gly Leu Asp Pro Asp Thr Asp Tyr Asp Val Ala Leu Val Pro
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Glu Ser Asn Val Arg Leu Leu Arg Pro Gln Ile
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<212> DNA
<213> Homo sapiens
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aataactata ctatcccaga agaagaaatt gggtctttct tatttcatgc tattaataag
120
ccaaatgete ctatctgget catactcaat gaagetggae tatactggag ageagtagga
aatagcaett ttgctattgc ctgtcttcag agggctttga atttagctcc acttcaatac
 caagatgttc ctcttgtcaa cttggccaac cttttgattc attacggcct tcatcttgat
 gccactaage tgctacttca agetttggcc atcaataget etgageetet gacetttttg
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 agcctgggaa atgcttacct tgctctgaag aatatcagtg gggcacttga ggcctttaga
 caggoettga aattaaccae caaatgteea gagtgtgaaa acageetgaa gttgateege
 tgtatgcagt tttatccttt tctgtacaac atcacttctt ctgtttgcag tggtaattgt
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 agcaatggtt ctgatgagat ggagaattca gatgaaacca aaatgtcaga agaaatactg
 gctttggtgg atgaatttca acaggcatgg cctttggaag gctttggggg tgcactagag
 840
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<210> 4692

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<212> PRT
<213> Homo sapiens
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Phe Leu Phe His Ala Ile Asn Lys Pro Asn Ala Pro Ile Trp Leu Ile
                           40
Leu Asn Glu Ala Gly Leu Tyr Trp Arg Ala Val Gly Asn Ser Thr Phe
Ala Ile Ala Cys Leu Gln Arg Ala Leu Asn Leu Ala Pro Leu Gln Tyr
Gln Asp Val Pro Leu Val Asn Leu Ala Asn Leu Leu Ile His Tyr Gly
Leu His Leu Asp Ala Thr Lys Leu Leu Cln Ala Leu Ala Ile Asn
                               105
Ser Ser Glu Pro Leu Thr Phe Leu Ser Leu Gly Asn Ala Tyr Leu Ala
Leu Lys Asn Ile Ser Gly Ala Leu Glu Ala Phe Arg Gln Ala Leu Lys
                       135
Leu Thr Thr Lys Cys Pro Glu Cys Glu Asn Ser Leu Lys Leu Ile Arg
                                       155
                   150
Cys Met Gln Phe Tyr Pro Phe Leu Tyr Asn Ile Thr Ser Ser Val Cys
                                   170
                165
Ser Gly Asn Cys His Glu Lys Thr Leu Asp Asn Ser His Asp Lys Gln
                               185
Lys Tyr Phe Asp Asn Ser Gln Ser Leu Asp Ala Ala Glu Glu Gro
                            200
Ser Glu Arg Gly Thr Glu Glu Asp Pro Val Phe Ser Val Glu Asn Ser
                        215
                                            220
Gly Arg Asp Ser Asp Ala Leu Arg Leu Glu Ser Thr Val Val Glu Glu
                    230
                                       235
Ser Asn Gly Ser Asp Glu Met Glu Asn Ser Asp Glu Thr Lys Met Ser
                245
                                    250
Glu Glu Ile Leu Ala Leu Val Asp Glu Phe Gln Gln Ala Trp Pro Leu
                                265
Glu Gly Phe Gly Gly Ala Leu Glu Met Lys Gly Arg Arg Leu Asp Leu
                           280
Gln Gly Ile Arg Val Leu Lys Lys Gly Pro Gln Asp Gly Val Ala Arg
                       295
Ser Ser Cys Tyr Gly Asp Cys Arg Ser Glu Asp Asp Glu Ala Thr Glu
                    310
                                        315
Trp Ile Thr Phe Gln Val Lys Arg Val Lys Lys Pro Lys Gly Asp His
               325
                                    330
Lys Lys Thr Pro Gly Lys Lys Val Glu Thr Gly Gln Ile Glu Asn Gly
                                345
His Arg Tyr Gln Ala Asn Leu Glu Ile Thr Gly Pro Lys Val Ala Ser
                           360
Pro Gly Pro Gln Gly Leu Leu Asp Trp Lys Thr Arg Lys Val Pro
                        375
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<211> 794
<212> DNA
<213> Homo sapiens
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<211> 103
<212> PRT
<213> Homo sapiens
<400> 4694
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                                 25
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Asn Ser Gly Val Gly Gln Asp Gly Ser Leu Leu Ser Ser Pro Phe Leu
        35
                             40
Lys Gly Phe Leu Ala Gly Tyr Val Val Ala Lys Leu Arg Ala Ser Ala
                         55
Val Leu Gly Phe Ala Val Gly Thr Cys Thr Gly Ile Tyr Ala Ala Gln
                    70
Ala Tyr Ala Val Pro Asn Val Glu Lys Thr Leu Arg Asp Tyr Leu Gln
                                     90
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Leu Leu Arg Lys Gly Pro Asp
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100

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1380

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160
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                    150
145
Lys Leu Ser Val Leu Leu Glu Lys Met Gly Gly Ser Ser Gly Ala
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                165
Leu Tyr Gly Leu Phe Leu Thr Ala Ala Ala Gln Pro Leu Lys Ala Lys
            180
                                185
Thr Ser Leu Pro Ala Trp Ser Ala Ala Met Asp Ala Gly Leu Glu Ala
                            200
                                                 205
Met Gln Lys Tyr Gly Lys Ala Ala Pro Gly Asp Arg Thr Met Leu Asp
                        215
Ser Leu Trp Ala Ala Glu Gln Glu Leu Gln Ala Trp Lys Ser Pro Gly
                                         235
                                                             240
                    230
225
Ala Asp Leu Leu Gln Val Leu Thr Lys Ala Val Lys Ser Ala Glu Ala
                                    250
                245
Ala Ala Glu Ala Thr Lys Asn Met Glu Ala Gly Ala Gly Arg Ala Ser
                                265
Tyr Ile Ser Ser Ala Arg Leu Glu Gln Pro Asp Pro Gly Ala Val Ala
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Ala Ala Ala Ile Leu Arg Ala Ile Leu Glu Val Leu Gln Ser
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 94n
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<212> PRT
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Thr Asp Gly Thr Val Phe Arg Ile His Thr Lys Ala Glu Gly Phe Met
                               25
Asp Ala Asp Ile Pro Leu Glu Leu Val Phe His Leu Pro Val Asn Tyr
                           40
Pro Ser Cys Leu Pro Gly Ile Ser Ile Asn Ser Glu Gln Leu Thr Arg
                                          60
Ala Gln Cys Val Thr Val Lys Glu Lys Leu Leu Glu Gln Ala Glu Ser
                   70
Leu Leu Ser Glu Pro Met Val His Glu Leu Val Leu Trp Ile Gln Gln
                                   90
Asn Leu Arg His Ile Leu Ser Gln Pro Glu Thr Gly Ser Gly Ser Glu
                               105
            100
Lys Cys Thr Phe Ser Thr Ser Thr Thr Met Asp Asp Gly Leu Trp Ile
                                              125
                           120
Thr Leu Leu His Leu Asp His Met Arg Ala Lys Thr Lys Tyr Val Lys
                       135
Ile Val Glu Lys Trp Ala Ser Asp Leu Arg Leu Thr Gly Arg Leu Met
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                   150
Phe Met Gly Lys Ile Ile Leu Ile Leu Leu Gln Gly Asp Arg Asn Asn
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Leu Lys Val Pro Lys Ser
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taagtcattt cacctcggag accgaaaaaa tgatcaaaaa gaaactatga gtaacaagct
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240
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gacttgatca gttaagaatt agttttcttg taaaacattc taaagccaag taaaatatcc
attottataa catacotata atatgagact aaggaatagg ttacatatag gtctacaaca
cattggtttg tctttaaaaa aacaaaagta gacatttata aataaaaaag agggacaatt
cacataggaa aaagaggtac acgagaaaat actgttgcac gcaataattt tcacacagat
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Lys His Thr Ala Phe Ala Thr Phe Pro Asn Glu Lys Ala Ala Ile Lys
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Glu Lys Pro Asn Lys Asp Leu Glu Ser Cys Ser Asp Asp Asp Asn Gln
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Cys Glu Gly Arg Thr Ala His Lys Ala Ala Arg Leu Gly Ile Thr Met
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Lys Ala Lys Leu Ala Arg Leu Glu Ala Gln Glu Gln Ala Phe Leu Ala
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Arg Leu Lys Gly Gln Asp Pro Gly Ala Pro Gln Leu Gln Ser Glu Ser
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Lys Pro Pro Lys Lys Lys Lys Lys Arg Arg Gln Lys Glu Glu Glu
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Glu Ala Thr Ala Ser Glu Arg Asn Asp Ala Asp Glu Lys His Pro Glu
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Arg Glu Glu Glu Glu Asn Asp Asp Asp Asn Ser Leu Glu Gly Glu
Thr Phe Pro Leu Glu Arg Asp Glu Val Met Pro Pro Pro Leu Gln His
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Pro Gln Thr Asp Arg Leu Thr Cys Pro Lys Gly Leu Pro Trp Ala Pro
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Lys Val Arg Glu Lys Asp Ile Glu Met Phe Leu Glu Ser Ser Arg Ser
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Lys Phe Ile Gly Tyr Thr Leu Gly Ser Asp Thr Asn Thr Val Val Gly
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Leu Pro Arg Pro Ile His Glu Ser Ile Lys Thr Leu Lys Gln His Lys
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                                         140
Tyr Thr Ser Ile Ala Glu Val Gln Ala Gln Met Lys Glu Glu Tyr Leu
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Arg Ser Pro Leu Ser Gly Gly Glu Glu Val Glu Gln Val Pro Ala
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                                                    175
Glu Thr Leu Tyr Gln Gly Leu Leu Pro Ser Leu Pro Gln Tyr Met Ile
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Ala Leu Leu Lys Ile Leu Leu Ala Ala Ala Pro Thr Ser Lys Ala Lys
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Thr Asp Ser Ile Asn Ile Leu Ala Asp Val Leu Pro Glu Glu Met Pro
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Thr Thr Val Leu Gln Ser Met Lys Leu Gly Val Asp Val Asn Arg His
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Asp Gly Gly Tyr Thr Ser Ser Cys Phe Asn Leu Ser Ala Met Phe Leu
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Gln Gly Ala Pro Gly Phe Pro Lys Asp Met Asp Leu Ala Cys Lys Tyr
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Ser Met Lys Ala Cys Asp Leu Gly His Ile Trp Ala Cys Ala Asn Ala
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Ser Arg Met Tyr
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                            40
Leu Thr Gly Glu Ser Glu Ser Ser Ser Glu Asp Glu Phe Glu Lys Glu
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Met Glu Ala Glu Leu Asn Ser Thr Met Lys Thr Met Glu Asp Lys Leu
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Ser Ser Leu Gly Thr Gly Ser Ser Ser Gly Asn Gly Lys Val Ala Thr
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Ala Pro Thr Arg Tyr Tyr Asp Asp Ile Tyr Phe Asp Ser Asp Ser Glu
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Asp Glu Asp Arg Ala Val Gln Val Thr Lys Lys Lys Lys Lys Gln
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His Lys Ile Pro Thr Asn Asp Glu Leu Leu Tyr Asp Pro Glu Lys Asp
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Asn Arg Asp Gln Ala Trp Val Asp Ala Gln Arg Arg Gly Tyr His Gly
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Ala Val Leu Asn Cys Pro Ala Cys Met Thr Thr Leu Cys Leu Asp Cys
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Gln Arg His Glu Ser Tyr Lys Thr Gln Tyr Arg Ala Met Phe Val Met
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                             200
Asn Cys Ser Ile Asn Lys Glu Glu Val Leu Arg Tyr Lys Ala Ser Glu
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Asn Arg Lys Lys Arg Arg Val His Lys Lys Met Arg Ser Asn Arg Glu
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                                         235
Asp Ala Ala Glu Lys Ala Glu Thr Asp Val Glu Glu Ile Tyr His Pro
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Gly Pro Pro Ser Pro Phe Pro Arg Gln Ser Pro Phe Gly Leu Asn Pro
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                            40
His Met Cys Thr Gly Ala Cys Ala Cys Val Asn Thr Cys Ser His Val
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Cys Thr Cys Xaa Ser Cys Pro Cys Xaa Tyr Val His Thr Cys Leu Cys
                                         75
Met His Ala Cys Ile Ala Val Cys Pro Tyr Pro His Val Arg Ile His
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 Val Gln Arg Phe Leu Tyr Thr Gly Arg Gln Pro Arg Pro Glu Glu Ala
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 Ala Glu Phe Gln Ala Glu Phe Val Ser Thr Pro Glu Leu Ala Ala Gln
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Leu Cys Asn Lys Asp Phe Phe Gln Lys Met Lys Glu Thr Ala Val Phe
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Ile Asn Ile Ser Arg Gly Asp Val Val Asn Gln Asp Asp Leu Tyr Gln
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                245
Ala Leu Ala Ser Gly Lys Ile Ala Ala Ala Gly Leu Asp Val Thr Ser
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Pro Glu Pro Leu Pro Thr Asn His Pro Leu Leu Thr Leu Lys Asn Cys
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                            280
Val Ile Leu Pro His Ile Gly Ser Ala Thr His Arg Thr Arg Asn Thr
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                                             300
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aatactggaa ttgctctttt tataattctc ttgacatttg tgtcaatatt ttccctgtat
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ggatataagg catctggatt agttggaaag ctt
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 <210> 4730
 <211> 148
 <212> PRT
 <213> Homo sapiens
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<400> 4730
Met Lys Lys Ala Glu Met Gly Arg Phe Ser Ile Ser Pro Asp Glu Asp
Ser Ser Ser Tyr Ser Ser Asn Ser Asp Phe Asn Tyr Ser Tyr Pro Thr
Lys Gln Ala Ala Leu Lys Ser His Tyr Ala Asp Val Asp Pro Glu Asn
Gln Asn Phe Leu Leu Glu Ser Asn Leu Gly Lys Lys Lys Tyr Glu Thr
                        55
Glu Phe His Pro Gly Thr Thr Ser Phe Gly Met Ser Val Phe Asn Leu
                    70
Ser Asn Ala Ile Val Gly Ser Gly Ile Leu Gly Leu Ser Tyr Ala Met
Ala Asn Thr Gly Ile Ala Leu Phe Ile Ile Leu Leu Thr Phe Val Ser
                                105
Ile Phe Ser Leu Tyr Ser Val His Leu Leu Leu Lys Thr Ala Asn Glu
        115
                            120
Gly Gly Ser Leu Leu Tyr Glu Gln Leu Gly Tyr Lys Ala Ser Gly Leu
                        135
    130
Val Gly Lys Leu
145
<210> 4731
<211> 2417
<212> DNA
<213> Homo sapiens
 <400> 4731
ttttttttt tttttcagac aaggtaccat tttattcctt ataaaatata tttcatattg
 ttgctgtaaa aacattacat ttcacatttt taaaaaaattt tttaacagta aaaataatac
 ttggaagaca gctgaggaaa aaggcgccaa taagacaaac tcacagatgg gatttatctc
 cctcttgctt ttttttttt tttttgcccc tggtaaaagt cagaacctgg gatgaccaga
 aagtaacagg acagatttct cccagcaaat cagtctccac aaccaaatga atattgttct
 ccaaggagtc aagctataga ctcacaatga caacgtggcc atggctcaaa acactctctg
 aaattacaaa attgctttct gagccaattt aaaagtcaca tgattgaatc caagctattt
 tactttaaat ggtccttttg ctttgcacct gagacctcgc ttggccacag acgtcattcg
 ctggactccc tgggcactaa atgagtgtct agcatcctta aggctgctca acacacagcc
 ccagactetg aatatgatte caagaaatat tetgaaaaaa gteacatege tggaataaac
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 gcaccttcct tccatcagag tctgctgccc gggtgggctg ggaaggaggg agatacaaag
 aagaaagtag gcatgatcac tgggtcggtt cccaagccac cctcaccctc caagaaggca
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tgaatggaac aaccccgaga acagagcacg tgtgaagaac caacacgaca ggcacgggat ggcagcactg gtggaaggga ggcaaggagg ccgccagtgc caaggaggag agggggcaca agogoaggoa gggaaggtgg caccaaaacc tagtaagaac aaagcaaaac caccgtggtt tocacactgo tototocott tattoctoto tttcctgccc tgtataccaa eggcataaga agectgcaca aagagaaaaa teegtatate cagttatate tacaeggtee aaactggggg cggqgggaat tcaaacagct ttctaaagac gagacggcag tgaaaactct gagggagagg 1140 atgaggacct gaagctgggg gttgtcttgg gaagtgaggg gggttgggaa acaccatcag cagetgecag etettaatte teaaggagat egaagggaea ggaaggagag eeetgegeea cctcaggcta gcctggcttt gagctttacc aagagacaga attccacata cattttttt tttttactaa gttataaaaa aaaaaacccc atcaccaaag acacctgtgc acaagtgtct gtocottotg toaccaacot agggoactac accottocca acateatgac cotactgoca 1500 ggtctacaga ttttgtaaca ctcaaagtgt cctgcattaa aaagcacgtg tctatttcct 1560 acgtgaaggg gccaagggag ccctggtggc ccaaatatct tcacccagga ctgggagggc 1620 ggcctcgatg acaaccaagg ggtggatgct gacactccat cccaggacag gtggctgggt 1680 aggatteeet gageeeetga eagetgggae atagggeeag gaettgtaee egaggeaget gggcagtggg cagtcacatt ccagtaggcc ctgaggaatc cccaaataag tcacgctggg 1800 aggaaagtga gacaccaaaa cagaaacatg ccctgccatc cgggcgtggc tcactctgtc ttcgcgcagg gctggttggc atggtgctac actcccgaga cctccctcct tctccccaag 1920 aacagetetg ettategaca tgcaegeage ecaggetece etagateeet ggaggeteca 1980 gaaacaccaa gggccaaaac gccagcagcc actaacccaa acccacgtot tootcotgto 2040 atttecteat cetgacgete aegggtgeaa ggacteteet tggeetteet eateetgett tcaggcagca aacagaaatg gggaaatccc tggtggggcc aggagacaga aaggaacctc 2160 cagaacctcc ctgggtctct cccggccacc caaataaaag aaaactttaa tcagtaaagg 2220 cttctgaata catcgtaaaa gaaaacaaag catttctgag gcgtcctttc aataaccgga 2280 ggaaggcggc gtcaggaggg tgcttcctcg ggtcagagca gagagtttcc agacgctcaa accetecagg agttectega ggaaagagga gagaatgate aaggtagtgt ttaactgeca 2400

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cattccaaaa agtgaat
2417
<210> 4732
<211> 129
<212> PRT
<213> Homo sapiens
<400> 4732
Met Ser Ile Ser Arg Ala Val Leu Gly Glu Lys Glu Gly Gly Leu Gly
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Ser Val Ala Pro Cys Gln Pro Ala Leu Arg Glu Asp Arg Val Ser His
                                                     30
Ala Arg Met Ala Gly His Val Ser Val Leu Val Ser His Phe Pro Pro
Ser Val Thr Tyr Leu Gly Ile Pro Gln Gly Leu Leu Glu Cys Asp Cys
                        55
Pro Leu Pro Ser Cys Leu Gly Tyr Lys Ser Trp Pro Tyr Val Pro Ala
                                         75
Val Arg Gly Ser Gly Asn Pro Thr Gln Pro Pro Val Leu Gly Trp Ser
                                     90
Val Ser Ile His Pro Leu Val Val Ile Glu Ala Ala Leu Pro Val Leu
                                 105
Gly Glu Asp Ile Trp Ala Thr Arg Ala Pro Leu Ala Pro Ser Arg Arg
                             120
                                                 125
        115
Lys
<210> 4733
<211> 543
<212> DNA
<213> Homo sapiens
<400> 4733
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ageggeegee gtgaccetet eggggateee aegatgttet tetacetgag caagaaaatt
tocattocca ataacgtgaa gotgoagtgt gtatootgga acaaggaaca agggttoata
gcatgcggtg gtgaagatgg attactgaaa gttttgaaat tagagacgca gacagatgat
 qcaaaattga ggggccttgc agcccccagt aacctttcta tgaatcagac tcttgaaggt
 catagtggtt ctgttcaagt tgtaacatgg aatgagcagt atcagaagtt gactaccagt
 gatgaaaacg ggcttatcat tgtgtggatg ttatataaag gctcttggat tgaggagatg
 420
 atcaacaatc gaaataaatc agttgttcgc agtatgagct ggaatgctga cggacagaag
 atotgoattg tatatgaaga tggggctgtg atagttggtt cagtggatgg caatogtatt
 540
 tqq
 543
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<210> 4734
<211> 181
<212> PRT
<213> Homo sapiens
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Xaa Pro Glu Leu Leu Val Leu Pro Ile Gly Asp Val Glu Pro Leu Leu
                                    10
Val Glu Gly Leu Ser Gly Arg Arg Asp Pro Leu Gly Asp Pro Thr Met
                                25
Phe Phe Tyr Leu Ser Lys Lys Ile Ser Ile Pro Asn Asn Val Lys Leu
        35
Gln Cys Val Ser Trp Asn Lys Glu Gln Gly Phe Ile Ala Cys Gly Gly
Glu Asp Gly Leu Leu Lys Val Leu Lys Leu Glu Thr Gln Thr Asp Asp
                                        75
Ala Lys Leu Arg Gly Leu Ala Ala Pro Ser Asn Leu Ser Met Asn Gln
                85
                                    90
Thr Leu Glu Gly His Ser Gly Ser Val Gln Val Val Thr Trp Asn Glu
                                                     110
                                 105
Gln Tyr Gln Lys Leu Thr Thr Ser Asp Glu Asn Gly Leu Ile Ile Val
                             120
                                                 125
Trp Met Leu Tyr Lys Gly Ser Trp Ile Glu Glu Met Ile Asn Asn Arg
                        135
                                             140
Asn Lys Ser Val Val Arg Ser Met Ser Trp Asn Ala Asp Gly Gln Lys
                                         155
                    150
145
Ile Cys Ile Val Tyr Glu Asp Gly Ala Val Ile Val Gly Ser Val Asp
                                                         175
                                     170
                165
Gly Asn Arg Ile Trp
            180
<210> 4735
<211> 300
 <212> DNA
 <213> Homo sapiens
<400> 4735
ntggtettet cagtacagea tggtggetgg ggcaggeega gagaatggea tggagaegee
gatgcacgag aacccggagt gggagaaggc ccgtcaggcc ctggccagca tcagcaagtc
aggagetgee ggeggetetg ecaagtecag cagcaatggg cetgtggeca gtgcacagta
 cgtgtcccag gcaaaagcct cagctttgca gcagcagcag tactaccagt ggtaccagca
 ggacaactat gcctacccct acagctacta ctatcccatg cccccaggcc ccggcatgga
 300
 <210> 4736
 <211> 93
 <212> PRT
 <213> Homo sapiens
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<400> 4736
Met Val Ala Gly Ala Gly Arg Glu Asn Gly Met Glu Thr Pro Met His
Glu Asn Pro Glu Trp Glu Lys Ala Arg Gln Ala Leu Ala Ser Ile Ser
                                25
Lys Ser Gly Ala Ala Gly Gly Ser Ala Lys Ser Ser Ser Asn Gly Pro
Val Ala Ser Ala Gln Tyr Val Ser Gln Ala Lys Ala Ser Ala Leu Gln
Gln Gln Gln Tyr Tyr Gln Trp Tyr Gln Gln Asp Asn Tyr Ala Tyr Pro
                    70
Tyr Ser Tyr Tyr Tyr Pro Met Pro Pro Gly Pro Gly Met
                85
<210> 4737
<211> 2602
<212> DNA
<213> Homo sapiens
<400> 4737
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aataatgtgg agatgtttcc accttcaggt tccactgggc tgattccccc ctcccacttt
 caagetegge ecettteaac tetgecaaga atggeteeca eetggetete agacatteee
 180
 ctggtccaac ccccaggcca tcaagatgtc tcagagaggc ggctagacac ccagagacct
 caagtgacca tgtgggaacg ggatgtttcc agtgacaggc aggagccagg gcggagaggc
 aggtcctggg ggctggaggg gtcacaggcc ctgagccagc aggctgaggt gatcgttcgg
 cagetgeaag agetgeggeg getggaggag gaggteegge teetgeggga gaeetegetg
 cagcagaaga tgaggctaga ggcccaggcc atggagctag aggctctggc acgggcggag
 aaggeeggee gagetgagge tgagggeetg egtgetgett tggetgggge tgaggttgte
 cggaagaact tggaagaggg gaggcagcgg gagctggaag aggttcagag gctgcaccaa
 gagcagctgt cctctttgac acaggctcac gaggaggctc tttccagttt gaccagcaag
 gctgagggct tggagaagtc tctgagtagt ctggaaacca gaagagcagg ggaagccaag
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gagetggeeg aggeteagag ggaggeegag etgettegga ageagetgag caagaeceag 780 gaagaettgg aggeteaggt gaceetggtt gagaatetaa gaaaatatt tggggaacaa 840 gteeettetg aggteeacag ceagacatgg gaactggage gacagaaget tetggaaace 900 atgeagetet tgeaggagga eegggacage etgeatgeea eegeggaget getgeaggtg 960

caacetteag attecetgga geetgagttt accaggaagt geeagteeet getgaacege tggcgggaga aggtgtttgc cctcatggtg cagctaaagg cccaggagct ggaacacagt gactctgtta agcagctgaa gggacaggtg gcctcactcc aggaaaaagt gacatcccag agccaggagc aggccatect geagegatee etgeaggaca aageegeaga ggtggaggtg gagcgtatgg gtgccaaggg cctgcagttg gagctgagcc gtgctcagga ggccaggcgt tggtggcagc agcagacagc ctcagccgag gagcagctga ggcttgtggt caatgctgtc 1380 agcagetete agatetgget egagaceace atggetaagg tggaagggge tgeegeecag 1440 cttcccagcc tcaacaaccg actcagctat gctgtccgca aggtccacac cattcggggc 1500 ctgattgctc gaaagcttgc ccttgctcag ctgcgccagg agagctgtcc cctaccacca ccggtcacag atgtgagcct tgagttgcag cagttgcggg aagaacggaa ccgcctggat gcagaactgc agctgagtgc ccgcctcatc cagcaggagg tgggccgggc tcgggagcaa 1680 ggggaggcag ageggeagea getgageaag gtggeecage agetggagea ggagetgeag cagacccagg agtccctggc tagcttgggg ctgcagctgg aggtagcacg ccagggccag caggagagca cagaggaggc tgccagtctg cggcaggagc tgacccagca gcaggaactc tacgggcaag ccctgcaaga aaaggtggct gaagtggaaa ctcggctgcg ggagcaactc tcagacacag agaggagget gaacgagget cggagggage atgccaagge cgtggtetee ttgcgccaga ttcagcgcag agccgcccag gaaaaggagc ggagccagga actcaggcgt ctgcaggagg aggcccggaa ggaggagggg cagcgactgg cccggcgctt gcaggagcta gagagggata agaacctcat gctggccacc ttgcagcagg aaggtctcct ctcccgttac aagcagcagc gactgttgac agttcttcct tccctactgg ataagaagaa atctgtggtg tecageecca ggeetecaga gtgttcagca tetgeacetg tagcageage agtgeecace agggagtcca taaaagggtc cctctctgtc ctgctcgatg acctgcagga cctgagtgaa gccatttcca aagaggaagc tgtttgtcaa ggagacaacc ttgacagatg ctccagctcc aatccccaga tgagcagcta agcagctgac agttggaggg aaagccagcc tgggggctgg gaggatectg gagaagtggg tggggacaga ccagceette eccatectgg ggttgeeetg ggggatacca gctgagtctg aattctgctc taaataaaga cgactacaga aggaaaaaaa 2580 aaaaaaaaaa aaaaaaaaaa aa 2602

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<210> 4738
<211> 756
<212> PRT
<213> Homo sapiens
<400> 4738
Met Ala Pro Thr Trp Leu Ser Asp Ile Pro Leu Val Gln Pro Pro Gly
His Gln Asp Val Ser Glu Arg Arg Leu Asp Thr Gln Arg Pro Gln Val
                                25
Thr Met Trp Glu Arg Asp Val Ser Ser Asp Arg Gln Glu Pro Gly Arg
                            40
Arg Gly Arg Ser Trp Gly Leu Glu Gly Ser Gln Ala Leu Ser Gln Gln
                        55
Ala Glu Val Ile Val Arg Gln Leu Gln Glu Leu Arg Arg Leu Glu Glu
                                        75
Glu Val Arg Leu Leu Arg Glu Thr Ser Leu Gln Gln Lys Met Arg Leu
                                    90
Glu Ala Gln Ala Met Glu Leu Glu Ala Leu Ala Arg Ala Glu Lys Ala
                                105
Gly Arg Ala Glu Ala Glu Gly Leu Arg Ala Ala Leu Ala Gly Ala Glu
                            120
Val Val Arg Lys Asn Leu Glu Glu Gly Arg Gln Arg Glu Leu Glu Glu
                       135
Val Gln Arg Leu His Gln Glu Gln Leu Ser Ser Leu Thr Gln Ala His
                                       155
                   150
Glu Glu Ala Leu Ser Ser Leu Thr Ser Lys Ala Glu Gly Leu Glu Lys
                                    170
                165
Ser Leu Ser Ser Leu Glu Thr Arg Arg Ala Gly Glu Ala Lys Glu Leu
                                185
Ala Glu Ala Gln Arg Glu Ala Glu Leu Leu Arg Lys Gln Leu Ser Lys
                            200
Thr Gln Glu Asp Leu Glu Ala Gln Val Thr Leu Val Glu Asn Leu Arg
                                            220
                        215
Lys Tyr Val Gly Glu Gln Val Pro Ser Glu Val His Ser Gln Thr Trp
                    230
                                        235
Glu Leu Glu Arg Gln Lys Leu Leu Glu Thr Met Gln Leu Leu Gln Glu
                                    250
                245
Asp Arg Asp Ser Leu His Ala Thr Ala Glu Leu Leu Gln Val Arg Val
                                265
            260
Gln Ser Leu Thr His Ile Leu Ala Leu Gln Glu Glu Glu Leu Thr Arg
                                                285
                            280
 Lys Val Gln Pro Ser Asp Ser Leu Glu Pro Glu Phe Thr Arg Lys Cys
                                            300
                        295
 Gln Ser Leu Leu Asn Arg Trp Arg Glu Lys Val Phe Ala Leu Met Val
                    310
                                        315
 Gln Leu Lys Ala Gln Glu Leu Glu His Ser Asp Ser Val Lys Gln Leu
                                     330
                325
 Lys Gly Gln Val Ala Ser Leu Gln Glu Lys Val Thr Ser Gln Ser Gln
                                345
             340
 Glu Gln Ala Ile Leu Gln Arg Ser Leu Gln Asp Lys Ala Ala Glu Val
                                                365
                             360
 Glu Val Glu Arg Met Gly Ala Lys Gly Leu Gln Leu Glu Leu Ser Arg
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375
    370
Ala Gln Glu Ala Arg Arg Trp Trp Gln Gln Gln Thr Ala Ser Ala Glu
                                       395
                   390
Glu Gln Leu Arg Leu Val Val Asn Ala Val Ser Ser Ser Gln Ile Trp
               405
                                    410
Leu Glu Thr Thr Met Ala Lys Val Glu Gly Ala Ala Ala Gln Leu Pro
                               425
Ser Leu Asn Asn Arg Leu Ser Tyr Ala Val Arg Lys Val His Thr Ile
                           440
                                               445
Arg Gly Leu Ile Ala Arg Lys Leu Ala Leu Ala Gln Leu Arg Gln Glu
                       455
Ser Cys Pro Leu Pro Pro Pro Val Thr Asp Val Ser Leu Glu Leu Gln
                                       475
                   470
Gln Leu Arg Glu Glu Arg Asn Arg Leu Asp Ala Glu Leu Gln Leu Ser
                                   490
                485
Ala Arg Leu Ile Gln Gln Glu Val Gly Arg Ala Arg Glu Gln Gly Glu
                                505
            500
Ala Glu Arg Gln Gln Leu Ser Lys Val Ala Gln Gln Leu Glu Gln Glu
                            520
Leu Gln Gln Thr Gln Glu Ser Leu Ala Ser Leu Gly Leu Gln Leu Glu
                        535
                                            540
Val Ala Arg Gln Gly Gln Gln Glu Ser Thr Glu Glu Ala Ala Ser Leu
                                       555
                    550
Arg Gln Glu Leu Thr Gln Gln Gln Glu Leu Tyr Gly Gln Ala Leu Gln
                                    570
                565
Glu Lys Val Ala Glu Val Glu Thr Arg Leu Arg Glu Gln Leu Ser Asp
                                585
Thr Glu Arg Arg Leu Asn Glu Ala Arg Arg Glu His Ala Lys Ala Val
                            600
Val Ser Leu Arg Gln Ile Gln Arg Arg Ala Ala Gln Glu Lys Glu Arg
                       615
Ser Gln Glu Leu Arg Arg Leu Gln Glu Glu Ala Arg Lys Glu Glu Gly
                    630
                                        635
Gln Arg Leu Ala Arg Arg Leu Gln Glu Leu Glu Arg Asp Lys Asn Leu
                                    650
Met Leu Ala Thr Leu Gln Gln Glu Gly Leu Leu Ser Arg Tyr Lys Gln
                                665
Gln Arg Leu Leu Thr Val Leu Pro Ser Leu Leu Asp Lys Lys Ser
                            680
Val Val Ser Ser Pro Arg Pro Pro Glu Cys Ser Ala Ser Ala Pro Val
                        695
Ala Ala Ala Val Pro Thr Arg Glu Ser Ile Lys Gly Ser Leu Ser Val
                    710
                                        715
Leu Leu Asp Asp Leu Gln Asp Leu Ser Glu Ala Ile Ser Lys Glu Glu
                                    730
Ala Val Cys Gln Gly Asp Asn Leu Asp Arg Cys Ser Ser Ser Asn Pro
                                745
            740
Gln Met Ser Ser
        755
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<210> 4739 <211> 684 <212> DNA <213> Homo sapiens

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tagccctctc tectgetect ttaaactetg aacttetagg atgggagaat gggaactttt
gcaggttgag attcatagtg aaatcgggtc aagaagtgat cagatgcaaa gcacagggca
gttcattact ataccatggc tgaggtcttc ctgggcacca ggccctgggc tcagcacttg
300
gctcagtctg caccttggac cctgccagag ccctccacag caggtgctct caggcaaggc
tgtgtgttgc tggccagacg ccttctgacc agcgtgcttt cttgaccaca gatcccttgg
ccaagcagga gggaaccatt agcagcctga ggagctggct ggctgggagc ctcggggacc
gcccagcett gctcccagct cacccacaag atgtggacag ctcttgtgct catttggatt
ttctccttgt ccttatctga aagccatgcg gcatccaacg atccacgtaa gtgagaaagc
tgtgtgactg ctggatgggc ccacggtggc cacaaagcat gctgagccct tgaaagcagc
atctgcaaac ccaqqccaac gcgt
684
<210> 4740
<211> 119
<212> PRT
<213> Homo sapiens
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Met Leu Leu Ser Arg Ala Gln His Ala Leu Trp Pro Pro Trp Ala His
Pro Ala Val Thr Gln Leu Ser His Leu Arg Gly Ser Leu Asp Ala Ala
Trp Leu Ser Asp Lys Asp Lys Glu Lys Ile Gln Met Ser Thr Arg Ala
Val His Ile Leu Trp Val Ser Trp Glu Gln Gly Trp Ala Val Pro Glu
                         55
Ala Pro Ser Gln Pro Ala Pro Gln Ala Ala Asn Gly Ser Leu Leu Leu
                                                             80
                     70
Gly Gln Gly Ile Cys Gly Gln Glu Ser Thr Leu Val Arg Arg Arg Leu
                                     90
Ala Ser Asn Thr Gln Pro Cys Leu Arg Ala Pro Ala Val Glu Gly Ser
            100
                                 105
Glv Arg Val Gln Gly Ala Asp
         115
<210> 4741
<211> 411
 <212> DNA
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<213> Homo sapiens

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ttttttctta aaaaaaaaa aggggttttt ctttgccccc cccgttcccc ccccttcccc
ttccgaaaaa aagaggggaa ttttttaaaa aacccgaaag gggggaaaggg ggggggtata
aaagataaaa tttggttttt tgggggggaa aatttggaca ccccaccctc gggttttttt
tccccacccc aaaaaatttt aaaagggggc cctaaaaaaa attttttctt taatttccaa
300
ataaaaaaa aatggggttc caaaatcatt gaaaaatagg ggggactcca aaaccttgaa
ttttcccaag ggggaccact aaaatttacc cettttttgg ggttttgggg g
411
<210> 4742
<211> 109
<212> PRT
<213> Homo sapiens
<400> 4742
Met Ile Leu Glu Pro His Phe Phe Phe Ile Trp Lys Leu Lys Lys Lys
Phe Phe Leu Gly Pro Pro Phe Lys Ile Phe Trp Gly Gly Glu Lys Lys
                                25
Pro Glu Gly Gly Val Ser Lys Phe Ser Pro Pro Lys Asn Gln Ile Leu
Ser Phe Ile Pro Pro Pro Phe Pro Pro Phe Gly Phe Phe Lys Lys Phe
                        55
Pro Ser Phe Phe Arg Lys Gly Lys Gly Gly Glu Arg Gly Gln Arg
                     70
                                        75
Lys Thr Pro Phe Phe Leu Arg Lys Lys Arg Glu Lys Lys Lys
                                     90
                 85
 Lys Glu Arg Lys Thr Pro Val Asp Leu Arg Glu Val Asn
             100
 <210> 4743
 <211> 473
 <212> DNA
 <213> Homo sapiens
 <400> 4743
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 caaccggccc cacaaattct agcagtgcca agaagaagga taaaagagtt caaggtggaa
 gagtgattga gtcccggtat ctgcagtatg aaaagaagac aacccaaaag gctcctgcag
 gagatgggtc acagacccga gggaagatgt ctgaaggtgg aaggaaatcc agcctgctcc
 agaaaagcaa agcagatagc agtggggtcg gaaagggtga cctgcagtcc acgttgctgg
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aagggcatgg cacageteca eetgaeetgg atetetetge tattaatgae aaaageateg
tcaaaaagac gccacagtta gcaaaaacaa tatcaaagaa acctgagtca acatcatttt
ctgcccctcg gaaaaagagc ccggatttat ctgaagcgaa tggaatgatg gag
473
<210> 4744
<211> 150
<212> PRT
<213> Homo sapiens
<400> 4744
Met Ala Asp Ser Ser Gly Arg Gly Ala Gly Lys Pro Ala Thr Gly Pro
Thr Asn Ser Ser Ser Ala Lys Lys Lys Asp Lys Arg Val Gln Gly Gly
            20
Arg Val Ile Glu Ser Arg Tyr Leu Gln Tyr Glu Lys Lys Thr Thr Gln
                             40
Lys Ala Pro Ala Gly Asp Gly Ser Gln Thr Arg Gly Lys Met Ser Glu
                                             60
Gly Gly Arg Lys Ser Ser Leu Leu Gln Lys Ser Lys Ala Asp Ser Ser
Gly Val Gly Lys Gly Asp Leu Gln Ser Thr Leu Leu Glu Gly His Gly
                                     90
 Thr Ala Pro Pro Asp Leu Asp Leu Ser Ala Ile Asn Asp Lys Ser Ile
                                 105
                                                     110
             100
 Val Lys Lys Thr Pro Gln Leu Ala Lys Thr Ile Ser Lys Lys Pro Glu
                                                  125
                             120
         115
 Ser Thr Ser Phe Ser Ala Pro Arg Lys Lys Ser Pro Asp Leu Ser Glu
                                              140
                         135
     130
 Ala Asn Gly Met Met Glu
                     150
 145
 <210> 4745
 <211> 666
 <212> DNA
 <213> Homo sapiens
 <400> 4745
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 gagcagctgg atcaccttgg tgaagttcag acggaatcag caggaattca gcgtgcacag
 attcagaaag aactttggcg aattcaggat gtcatggaag ggctgagtaa acataagcag
 caaagaggta ctacagaaat aggtatgata ggatcaaagc ctttctcaac agttaagtac
  aaaaatgagg gtccagatta tagactctac aagagtgaac cagagttaac aacagtggca
  gaagttgatg aatctaatgg agaagaaaaa tcagaacctg tttcagagat agaaacttca
  gttgttaaag gttcccactt tcctgttgga gtagtccctc caagagcaaa atcaccaaca
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Gln Asp Val Met Glu Gly Leu Ser Lys His Lys Gln Gln Arg Gly Thr
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Thr Glu Ile Gly Met Ile Gly Ser Lys Pro Phe Ser Thr Val Lys Tyr
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Lys Asn Glu Gly Pro Asp Tyr Arg Leu Tyr Lys Ser Glu Pro Glu Leu
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Cys Glu Gln Asn Leu Leu Ser Arg Pro Asp Gly Ser Ala Ser Phe Leu
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Gln Gly Asp Thr Ser Val Leu Ala Gly Val Tyr Gly Pro Ala Glu Val
Lys Val Ser Lys Glu Ile Phe Asn Lys Ala Thr Leu Glu Val Ile Leu
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Glu Lys Ser Arg Glu Arg Leu Ile Arg Asn Thr Cys Glu Ala Val Val
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Leu Gly Thr Leu His Pro Arg Thr Ser Ile Thr Val Val Leu Gln Val
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Val Ser Asp Ala Gly Ser Leu Leu Ala Cys Cys Leu Asn Ala Ala Cys
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Met Ala Leu Val Asp Ala Gly Val Pro Met Arg Ala Leu Phe Cys Gly
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Val Ala Cys Ala Leu Asp Ser Asp Gly Thr Leu Val Leu Asp Pro Thr
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Ser Lys Gln Glu Lys Glu Ala Arg Ala Val Leu Thr Phe Ala Leu Asp
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Ser Val Glu Arg Lys Leu Leu Met Ser Ser Thr Lys Gly Leu Tyr Ser
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Asp Thr Glu Leu Gln Gln Cys Leu Ala Ala Ala Gln Ala Ala Ser Gln
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Thr Gln Thr Ser Ile Thr Asp Ser Cys Ala Val Tyr Arg Val Asn Asn
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Val Phe Val Val Asp Ser Ala Ala Phe Gln Arg Glu Val Lys Asp Val
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Lys Ser Ala Lys Leu Ile Gln Gln Gln Leu Glu Lys Glu Leu Asn Thr
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Leu Arg Val Thr Arg Ser Ala Ala Pro Ser Thr Leu Asp Ser Ser Ser
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Lys Leu Ala Phe Met Val Ser Leu Gly Leu Val Thr His Asp His Leu
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Ala Asn Glu Glu His Trp Pro Lys Gly Asp Ile His Glu Asp Phe Cys
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Ser Val Cys Arg Lys Ser Gly Gln Leu Leu Met Cys Asp Thr Cys Ser
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Arg Val Tyr His Leu Asp Cys Leu Asp Pro Pro Leu Lys Thr Ile Pro
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Lys Gly Met Trp Ile Cys Pro Arg Cys Gln Asp Gln Met Leu Lys Lys
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Ser Asp Val Glu Gly Gly Glu Val Leu Tyr Leu Val His Tyr Cys Gly
Trp Asn Val Arg Tyr Asp Glu Trp Ile Lys Ala Asp Lys Ile Val Arg
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Pro Ala Asp Lys Asn Val Pro Lys Ile Lys His Arg Lys Lys Ile Lys
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Asn Lys Leu Asp Lys Glu Lys Asp Lys Asp Glu Lys Tyr Ser Pro Lys
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Asn Cys Lys Leu Arg Arg Leu Ser Lys Pro Pro Phe Gln Thr Asn Pro
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Ser Pro Glu Met Val Ser Lys Leu Asp Leu Thr Asp Ala Lys Asn Ser
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Asp Thr Ala His Ile Lys Ser Ile Glu Ile Thr Ser Ile Leu Asn Gly
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Leu Gln Ala Ser Glu Ser Ser Ala Glu Asp Ser Glu Gln Glu Asp Glu
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Asp His Leu Thr Asn Asn Arg Asn Asp Leu Ile Ser Lys Glu Glu Gln
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Gln Ile Lys Arg Gly Lys Arg Arg Tyr Cys Asn Thr Glu Glu Cys Leu
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Phe Asp Ala Ser Val Ser Ser Ser Ser Ser Asn Gln Pro Glu Pro Glu
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Val Val Asn Asn Thr Lys Lys Gly Lys Gly Thr Asn Ser Ser Asp Ser
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Ser Val Ser Thr Gly Met Lys Ser His Ser Thr Lys Ser Pro Ala Arg
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Thr Gln Ser Pro Gly Lys Cys Gly Lys Asn Gly Asp Lys Asp Pro Asp
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Gln Met Ser Asp Leu Glu Asn Met Thr Ser Ala Glu Arg Ile Thr Ile
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Leu Gln Glu Lys Leu Gln Glu Asn Gln Lys His Tyr Leu Ser Leu Lys
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Ser Glu Val Ala Ser Ile Asp Arg Arg Arg Lys Arg Leu Lys Lys Lys
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Glu Gln Arg Arg Pro Ser Thr Ser Ser Ala Ser Gly Gln Trp Ser Pro
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Thr Pro Glu Trp Val Leu Ser Trp Lys Ser Lys Leu Pro Leu Gln Thr
Ile Met Arg Leu Leu Gln Val Leu Val Pro Gln Val Glu Lys Ile Cys
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Ile Asp Lys Gly Leu Thr Asp Glu Ser Glu Ile Leu Arg Phe Leu Gln
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Arg Lys Tyr Gln Ala Asn Ser Gly Thr Ala Met Trp Phe Arg Thr Tyr
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Met Trp Gly Val Ile Tyr Leu Arg Asn Val Asp Pro Pro Val Trp Tyr
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Lys Gly Gln Thr Lys Thr Leu Phe Glu Phe Ser Ser Arg Ala Gly
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Arg Ala Ser Gly Tyr Arg Lys Arg Gly Pro Lys Pro Lys Arg Leu Leu
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Leu Gln Arg Leu Tyr Ser Met Asp Leu Arg Ser Ser His Lys Ala Lys
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Gly Lys Glu Lys Leu Cys Phe Ser Leu Thr Cys Pro Leu Gly Ser Gly
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Ser Pro Glu Gly Val Val Lys Ala Gly Ala Pro Glu Leu Val Asp Lys
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Gly Pro Leu Val Pro Thr Leu Pro Phe Pro Leu Arg Lys Pro Arg Lys
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Ala His Lys Tyr Leu Arg Leu Ser Arg Lys Lys Phe Pro Pro Arg Gly
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aggtcaacaa gaacaaggaa atgtataagc tacagatttg atgagtttga tgaagcaatt
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gatgaagcta ttgaagatga catcaaagaa gccgatggag gaggagttgg ccgaggaaaa
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Leu Lys Lys Arg Glu Ile Lys Leu Ser Asp Asp Phe Asp Ser Pro Val
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Lys Gly Pro Leu Cys Lys Ser Val Thr Pro Thr Lys Glu Phe Leu Lys
Asp Glu Ile Lys Gln Glu Glu Glu Thr Cys Lys Arg Ile Ser Thr Ile
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 Thr Ala Leu Gly His Glu Gly Lys Gln Leu Val Asn Gly Glu Val Ser
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 Asp Glu Arg Val Ala Pro Asn Phe Lys Thr Glu Pro Ile Glu Thr Lys
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 Phe Tyr Glu Thr Lys Glu Glu Ser Tyr Ser Pro Ser Lys Asp Arg Asn
                                                 125
                             120
 Ile Ile Thr Glu Gly Asn Gly Thr Glu Ser Leu Asn Ser Val Ile Thr
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                                             140
 Ser Met Lys Thr Gly Glu Leu Glu Lys Glu Thr Ala Pro Leu Arg Lys
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 Asp Ala Asp Ser Ser Ile Ser Val Leu Glu Ile His Ser Gln Lys Ala
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 Gln Ile Glu Glu Pro Asp Pro Pro Glu Met Glu Thr Ser Leu Asp Ser
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Ser Glu Met Ala Lys Asp Leu Ser Ser Lys Thr Ala Leu Ser Ser Thr
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Glu Ser Cys Thr Met Lys Gly Glu Glu Lys Ser Pro Lys Thr Lys Lys
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Asp Lys Arg Pro Pro Ile Leu Glu Cys Leu Glu Lys Leu Glu Lys Ser
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Lys Lys Thr Phe Leu Asp Lys Asp Ala Gln Arg Leu Ser Pro Ile Pro
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Glu Glu Val Pro Lys Ser Thr Leu Glu Ser Glu Lys Pro Gly Ser Pro
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                               265
Glu Ala Ala Glu Thr Ser Pro Pro Ser Asn Ile Ile Asp His Cys Glu
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Lys Leu Ala Ser Glu Lys Glu Val Val Glu Cys Gln Ser Thr Ser Thr
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Val Gly Gly Gln Ser Val Lys Lys Val Asp Leu Glu Thr Leu Lys Glu
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Asp Ser Glu Phe Thr Lys Val Glu Met Asp Asn Leu Asp Asn Ala Gln
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Thr Ser Gly Ile Glu Glu Pro Ser Glu Thr Lys Gly Ser Met Gln Lys
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Ser Lys Phe Lys Tyr Lys Leu Val Pro Glu Glu Glu Thr Thr Ala Ser
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Glu Asn Thr Glu Ile Thr Ser Glu Arg Gln Lys Glu Gly Ile Lys Leu
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                                           380
Thr Ile Arg Ile Ser Ser Arg Lys Lys Lys Pro Asp Ser Pro Pro Lys
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Val Leu Glu Pro Glu Asn Lys Gln Glu Lys Thr Glu Lys Glu Glu Glu
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Lys Thr Asn Val Gly Arg Thr Leu Arg Arg Ser Pro Arg Ile Ser Arg
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Pro Thr Ala Lys Val Ala Glu Ile Arg Asp Gln Lys Ala Asp Lys Lys
                            440
Arq Gly Glu Gly Glu Asp Glu Val Glu Glu Glu Ser Thr Ala Leu Gln
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Lys Thr Asp Lys Lys Glu Ile Leu Lys Lys Ser Glu Lys Asp Thr Asn
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Ser Lys Val Ser Lys Val Lys Pro Xaa Lys Ala Lys Phe Asp Gly Leu
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Val Leu Gly His Val Ala Asp Gly Asn Ile Pro Ala Met Met Lys Val
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Lys Gly Leu Ala Val Lys Asn His Leu Gln Leu Gln Lys Arg Arg Lys
Lys Arg Lys Val Lys Lys Pro Ser Xaa Ala Asp Asp Asp Glu Pro Cys
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Lys Lys Cys Gly Leu Pro Asn His Pro Glu Leu Ile Leu Leu Cys Asp
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Ser Cys Asp Ser Gly Tyr His Thr Ala Cys Leu Arg Pro Pro Leu Met
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                565
 Ile Ile Pro Asp Gly Glu Trp Phe Cys Pro Pro Cys Gln His Lys Leu
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 Leu Cys Glu Lys Leu Glu Glu Gln Leu Gln Asp Leu Asp Val Ala Leu
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 Lys Lys Lys Glu Arg Ala Glu Arg Arg Lys Glu Arg Leu Val Tyr Val
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620
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Gly Ile Ser Ile Glu Asn Ile Ile Pro Pro Gln Glu Pro Asp Phe Ser
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625
Glu Asp Gln Glu Glu Lys Lys Lys Asp Ser Lys Lys Ser Lys Ala Asn
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Leu Leu Glu Arg Arg Ser Thr Arg Thr Arg Lys Cys Ile Ser Tyr Arg
Phe Asp Glu Phe Asp Glu Ala Ile Asp Glu Ala Ile Glu Asp Asp Ile
                            680
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Lys Glu Ala Asp Gly Gly Gly Val Gly Arg Gly Lys Asp Ile Ser Thr
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Ile Thr Gly His Arg Gly Lys Asp Ile Ser Thr Ile Leu Asp Glu
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  atgggttctc gtgtttcctg ttcaggatca ccagcatttc tgagcttggg tttatgcacg
  1020
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tatttaacag tcacaagaag tcttatttac atgccaccaa ccaacctcag aaacccataa
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Pro Glu Pro Arg Arg Thr Glu His Arg Ala Pro Ser Ser Thr Trp Arg
                            40
Pro Val Ala Leu Thr Leu Leu Thr Leu Cys Leu Val Leu Leu Ile Gly
                        55
Leu Ala Ala Leu Gly Leu Leu Phe Phe Gln Tyr Tyr Gln Leu Ser Asn
                    70
                                         75
Thr Gly Gln Asp Thr Ile Ser Gln Met Glu Glu Arg Leu Gly Asn Thr
                                     90
                85
Ser Gln Glu Leu Gln Ser Leu Gln Val Gln Asn Ile Lys Leu Ala Gly
            100
                                105
Ser Leu Gln His Val Ala Glu Lys Leu Cys Arg Glu Leu Tyr Asn Lys
                            120
Ala Gly Ala His Arg Cys Ser Pro Cys Thr Glu Gln Trp Lys Trp His
                        135
                                             140
Gly Asp Asn Cys Tyr Gln Phe Tyr Lys Asp Ser Lys Ser Trp Glu Asp
                    150
                                         155
145
Cys Lys Tyr Phe Cys Leu Ser Glu Asn Ser Thr Met Leu Lys Ile Asn
                                     170
Lys Gln Glu Asp Leu Glu Phe Ala Ala Ser Gln Ser Tyr Ser Glu Phe
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Phe Tyr Ser Tyr Trp Thr Gly Leu Leu Arg Pro Asp Ser Gly Lys Ala
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Trp Leu Trp Met Asp Gly Thr Pro Phe Thr Ser Glu Leu Phe His Ile
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Ile Ile Asp Val Thr Ser Pro Arg Ser Arg Asp Cys Val Ala Ile Leu
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                                        235
Asn Gly Met Ile Phe Ser Lys Asp Cys Lys Glu Leu Lys Arg Cys Val
                                    250
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Cys Glu Arg Arg Ala Gly Met Val Lys Pro Glu Ser Leu His Val Pro
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Pro Glu Thr Leu Gly Glu Gly Asp
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gaggatgcac tgtcccctgt ggacatggag cccgagaagc tggtgcacaa gttcaaggag
960
ctccagatca agcatgcggt cactgaggca gagatccagc agctgaaaag aaagctgcag
1020
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Asp Phe Ser Glu Ala Asp Leu Val Asp Val Ser Ala Tyr Ser Gly Leu
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Gly Glu Asp Ser Ala Gly Ser Ala Leu Glu Glu Asp Asp Glu Asp Asp
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Glu Gly Asp Gly Glu Pro Pro Tyr Glu Pro Glu Ser Gly Cys Val Glu
                                        75
                    70
Ile Pro Gly Leu Ser Glu Glu Glu Asp Pro Ala Pro Ser Arg Lys Ile
                                    90
His Phe Ser Thr Ala Pro Ile Gln Val Phe Ser Thr Tyr Ser Asn Glu
                                105
Asp Tyr Asp Arg Arg Asn Glu Asp Val Asp Pro Met Ala Ala Ser Ala
                                                125
                            120
Glu Tyr Glu Leu Glu Lys Arg Val Glu Arg Leu Glu Leu Phe Pro Val
                        135
                                             140
Glu Leu Glu Lys Asp Ser Glu Gly Leu Gly Ile Ser Ile Ile Gly Met
                    150
                                         155
Gly Ala Gly Ala Asp Met Gly Leu Glu Lys Leu Gly Ile Phe Val Lys
                                     170
Thr Val Thr Glu Gly Gly Ala Ala His Arg Asp Gly Arg Ile Gln Val
                                                     190
                                 185
Asn Asp Leu Leu Val Glu Val Asp Gly Thr Ser Leu Val Gly Val Thr
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Gln Ser Phe Ala Ala Ser Val Leu Arg Asn Thr Lys Gly Arg Val Arg
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                                             220
Phe Met Ile Gly Arg Glu Arg Pro Gly Glu Gln Ser Glu Val Ala Gln
                                         235
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Leu Ile Gln Gln Thr Leu Glu Gln Glu Arg Trp Gln Arg Glu Met Met
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Glu Gln Arg Tyr Ala Gln Tyr Gly Glu Asp Asp Glu Glu Thr Gly Glu
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                                 265
 Tyr Ala Thr Asp Glu Asp Glu Glu Leu Ser Pro Thr Phe Pro Gly Gly
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Glu Met Ala Ile Glu Val Phe Glu Leu Ala Glu Asn Glu Asp Ala Leu
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Ser Pro Val Asp Met Glu Pro Glu Lys Leu Val His Lys Phe Lys Glu
305
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Leu Gln Ile Lys His Ala Val Thr Glu Ala Glu Ile Gln Gln Leu Lys
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                                   330
               325
Arg Lys Leu Gln Ser Leu Glu Gln Glu Lys Gly Arg Trp Arg Val Glu
                               345
Lys Ala Gln Leu Glu Gln Ser Val Glu Glu Asn Lys Glu Arg Met Glu
        355
                           360
Lys Leu Glu Gly Tyr Trp Gly Glu Ala Gln Ser Leu Cys Gln Ala Val
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Asp Glu His Leu Arg Glu Thr Gln Ala Gln Tyr Gln Ala Leu Glu Arg
                                       395
                                                           400
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Lys Tyr Ser Lys Ala Lys Arg Leu Ile Lys Asp Tyr Gln Gln Lys Glu
                                   410
Ile Glu Phe Leu Lys Lys Glu Thr Ala Gln Arg Arg Val Leu Glu Glu
            420
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Ser Glu Leu Ala Arg Lys Glu Glu Met Asp Lys Leu Leu Asp Lys Ile
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Ser Glu Leu Glu Gly Asn Leu Gln Thr Leu Arg Asn
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caacagagca tacatacctt ggaagggtgt gttctgatat actcgtatgg aaagttctga
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600
caagaagacc gttgcagtcc agagatgaga aactggacca gaggcaaatc atgaacagaa
cgggagtcaa gagaaggggt ttctaagatg gagaagtggg ggcgggtgtg gatccagtgg
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Leu Ser Val Leu Thr Glu Cys Ala Arg Met His Arg Pro Ala Arg Lys
Phe Leu Lys Ala Gln Val Leu Pro Pro Leu Arg Asp Val Arg Thr Arg
Pro Glu Val Gly Asp Leu Leu Arg Asn Lys Leu Val Arg Leu Met Thr
                                                             80
                    70
His Leu Asp Thr Asp Val Lys Arg Val Ala Ala Glu Phe Leu Phe Val
                                     90
Leu Cys Ser Glu Ser Val Pro Arg Phe Ile Lys Tyr Thr Gly Tyr Gly
                                 105
Asn Ala Ala Gly Leu Leu Ala Ala Arg Gly Leu Met Ala Gly Gly Arg
        115
 Pro Glu Gly Gln Tyr Ser Glu Asp Glu Asp Thr Asp Thr Asp Glu Tyr
     130
                         135
 Lys Glu Ala Lys Ala Ser Ile Asn Pro Val Thr Gly Arg Val Glu Glu
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Lys Pro Pro Asn Pro Met Glu Gly Met Thr Glu Glu Gln Lys Glu His
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Glu Ala Met Lys Leu Val Thr Met Phe Asp Lys Leu Ser Ser Pro Thr
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Ala Pro Phe Pro Asn Arg Asn Arg Val Ile Gln Pro Met Gly Met Ser
                            200
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Pro Arg Gly His Leu Thr Ser Leu Gln Asp Ala Met Cys Glu Thr Met
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Glu Gln Gln Leu Ser Ser Asp Pro Asp Ser Asp Pro Asp
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<210> 4771
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 cccaggetag accetecata cacacecagg ctagacecte catacteace caggetagae
 1080
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Lys Pro Asp Val Val Gln Asp Lys Glu Thr Glu Arg Asn Leu Gln Arg
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Ile Ala Thr Arg Gly Val Val Gln Leu Phe Asn Ala Val Gln Lys His
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Gln Lys Asn Val Asp Glu Lys Val Lys Glu Ala Gly Ser Ser Met Arg
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Lys Arg Ala Lys Leu Ile Ser Thr Val Ser Lys Lys Asp Phe Ile Ser
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Val Leu Arg Gly Met Asp Gly Ser Thr Asn Glu Thr Ala Ser Ser Arg
                            120
Lys Lys Pro Lys Ala Lys Gln Thr Glu Val Lys Ser Glu Glu Gly Pro
                        135
Gly Trp Thr Ile Leu Arg Asp Asp Phe Met Met Gly Ala Ser Met Lys
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Asp Trp Asp Lys Glu Ser Asp Gly Pro Asp Asp Ser Arg Pro Glu Ser
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Ala Ser Asp Ser Asp Thr
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Ile His Ala Ala His Pro Val Thr Ser Phe Gln Phe Leu Leu Thr Phe

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Lys Asp Ile Arg Glu Asn Phe Gln Asp Leu Met Gln Tyr Cys Ala Gln
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Sei	r Phe	e Tr	o Arg	Ası	a Ala	a His	s Lys	arç	; Ile	e Sea	: se:	Glr	1 Met	. Val	Val
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Cys Ser Asn Ile Thr Asn Thr Gly Leu Leu Leu Ile Ala Trp Gly Leu
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Cys Leu Gly Leu Glu Gln Leu Thr Leu Gln Asp Cys Gln Lys Leu Thr
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Pro His Ser Ile Ile Asn Gly Tyr Lys Arg Val Leu Ala Leu Ser Val
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Gly Pro Val Ile Ile Pro Gln Met Leu Leu Glu Leu Trp Ala Gln Gly
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<213> Homo sapiens
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Pro Pro Cys Gly His Arg Gly Ala Leu Asp Gln Pro His His Arg Val
                            40
Ala Gln Pro His Leu Gln Val Val Arg Gln Arg Ser Pro Pro Ala Ser
                                             60
                        55
Trp Ser Pro Pro Pro Arg Ala Leu Ser His Val Phe Leu Phe Gly Asp
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Arg Pro Phe Trp Trp Val His Glu Ser Gly Tyr Tyr Ser Gln Ala Pro
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Ala Gln Val His Gln Phe Pro Ser Ser Cys Glu Thr Gly Pro Gly Ser
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Pro Ser Gly His Cys Met Ile
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1140
gtgtgcctcc ctttcctttc cctcccacaa agccaacact ctgtgaccac cacactccag
gaggeagece cateccette cagecectaa gtaggeeete eceteectaa atetgettee
1260
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gcaccacetg gtcttagccc caaagatggg ccttctctct cccagataag ttggtcctcc
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Arg Pro Gly Ala Ser Arg Gly Leu Val Gly Ser Trp Ala Ala Met Glu
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Ser Thr Leu Gly Ala Gly Ile Val Ile Ala Glu Ala Leu Gln Asn Gln
                            40
Leu Ala Trp Leu Glu Asn Val Trp Leu Trp Ile Thr Phe Leu Gly Asp
                        55
Pro Lys Ile Leu Phe Leu Phe Tyr Phe Pro Ala Ala Tyr Tyr Ala Ser
                                        75
                    70
Arg Arg Val Gly Ile Ala Val Leu Trp Ile Ser Leu Ile Thr Glu Trp
                                    90
                85
Leu Asn Leu Ile Phe Lys Trp Phe Leu Phe Gly Asp Arg Pro Phe Trp
                                105
Trp Val His Glu Ser Gly Tyr Tyr Ser Gln Ala Pro Ala Gln Val His
                            120
Gln Phe Pro Ser Ser Cys Glu Thr Gly Pro Gly Ser Pro Ser Gly His
                        135
                                             140
Cys Met Ile Thr Gly Ala Ala Leu Trp Pro Ile Met Thr Ala Leu Ser
                                        155
                    150
Ser Gln Val Ala Thr Arg Ala Arg Ser Arg Trp Val Arg Val Met Pro
                                    170
Ser Leu Ala Tyr Cys Thr Phe Leu Leu Ala Val Gly Leu Ser Arg Ile
                                185
Phe Ile Leu Ala His Phe Pro His Gln Val Leu Ala Gly Leu Ile Thr
                            200
                                                 205
Gly Ala Val Leu Gly Trp Leu Met Thr Xaa Pro Glu Cys Leu Trp Ser
                                             220
                         215
Gly Ser Xaa Ser Phe Tyr Gly Leu Thr Ala Leu Ala Leu Met Leu Gly
                                         235
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Thr Ser Leu Ile Tyr Trp Thr Leu Phe Thr Leu Gly Leu Asp Leu Ser
                                    250
                245
Trp Ser Ile Ser Leu Ala Phe Lys Trp Cys Glu Arg Pro Glu Trp Ile
                                265
His Val Asp Ser Arg Pro Phe Ala Ser Leu Ser Arg Asp Ser Gly Ala
                                                 285
 Ala Leu Gly Leu Gly Ile Ala Leu His Ser Pro Cys Tyr Ala Gln Val
                                             300
 Arg Arg Ala Gln Leu Gly Asn Gly Gln Lys Ile Ala Cys Leu Val Leu
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320
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305
Ala Met Gly Leu Leu Gly Pro Leu Asp Trp Leu Gly His Pro Pro Gln
                                    330
                325
Ile Ser Leu Phe Tyr Ile Phe Asn Phe Leu Lys Tyr Thr Leu Trp Pro
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Cys Leu Val Leu Ala Leu Val Pro Trp Ala Val His Met Phe Ser Ala
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Gln Glu Ala Pro Pro Ile His Ser Ser
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<212> DNA
<213> Homo sapiens
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ccaaaacctg ctaatgcctg atttccatta cgtgctactc ctcaaatggc agcggcttct
gaatattaca gagatggtgt gctgtttgct tttctctttt gttgtagcat aaaactgttc
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qaaqttaaat ttagatgett teetetette ttattttgtg gaggtattte etgtteagta
qcaaatcagt tatagaatat attagcattg ttatatttta aactaatgac taatcatttc
agotttatto atactgttgo attttatatt toacagggag caatagaaaa agtgaaagaa
agtgacaaac tagttgcaac aagtaaaatc accctacaag acaaacagaa catggtgaag
agagtcagca tcatgtctta cgcg
564
<210> 4804
<211> 53
<212> PRT
<213> Homo sapiens
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Gln Gly Ala Ile Glu Lys Val Lys Glu Ser Asp Lys Leu Val Ala Thr
                                 25
 Ser Lys Ile Thr Leu Gln Asp Lys Gln Asn Met Val Lys Arg Val Ser
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 Ile Met Ser Tyr Ala
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<212> DNA <400> 4805

<213> Homo sapiens

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aatactcaga taggtataag atttttcaca aaatccttat gtaagataca ttccattttt
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<211> 438
<212> PRT
<213> Homo sapiens
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Glu Ile Phe Thr Thr Asn Asn Gly Tyr Lys Ser Met Gln Lys Lys Leu
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Arg Ser Asn Trp Lys Ile Gln Ser Leu Lys Asp Glu Ile Thr Ser Glu
Lys Leu Asn Gly Val Lys Leu Trp Ile Thr Ala Gly Pro Arg Glu Lys
                       55
                                           60
Phe Thr Ala Ala Glu Phe Glu Ile Leu Lys Lys Tyr Leu Asp Thr Gly
Gly Asp Val Leu Val Met Leu Gly Glu Gly Glu Ser Arg Phe Asp
               85
Thr Asn Ile Asn Phe Leu Leu Glu Glu Tyr Gly Ile Met Val Asn Asn
                               105
Asp Ala Val Val Arg Asn Val Tyr His Lys Tyr Phe His Pro Lys Glu
                                              125
                           120
Ala Leu Val Ser Ser Gly Val Leu Asn Arg Glu Ile Ser Arg Ala Ala
                                           140
                       135
Gly Lys Ala Val Leu Ala Ile Ile Asp Glu Glu Ser Ser Gly Asn Asn
                                       155
                    150
Ala Gln Ala Leu Thr Phe Val Tyr Pro Phe Gly Ala Thr Leu Ser Val
                165
                                   170
Met Lys Pro Ala Val Ala Val Leu Ser Thr Gly Ser Val Cys Phe Pro
                               185
                                                   190
Leu Asn Arg Pro Ile Leu Ala Phe Tyr His Ser Lys Asn Gln Gly Gly
                                               205
                            200
Lys Leu Ala Val Leu Gly Ser Cys His Met Phe Ser Asp Gln Tyr Leu
                                           220
                       215
 Asp Lys Glu Glu Asn Ser Lys Ile Met Asp Val Val Phe Gln Trp
                    230
                                       235
 Leu Thr Thr Gly Asp Ile His Leu Asn Gln Ile Asp Ala Glu Asp Pro
                                   250
                245
 Glu Ile Ser Asp Tyr Met Met Leu Pro Tyr Thr Ala Thr Leu Ser Lys
                                265
 Arg Asn Arg Glu Cys Leu Gln Glu Ser Asp Glu Ile Pro Arg Asp Phe
                            280
 Thr Thr Leu Phe Asp Leu Ser Ile Phe Gln Leu Asp Thr Thr Ser Phe
                                           300
                        295
 His Ser Val Ile Glu Ala His Glu Gln Leu Asn Val Lys His Glu Pro
                                       315
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 Leu Gln Leu Ile Gln Pro Gln Phe Glu Thr Pro Leu Pro Thr Leu Gln
                                   330
 Pro Ala Val Phe Pro Pro Ser Phe Arg Glu Leu Pro Pro Pro Pro Leu
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350
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Glu Leu Phe Asp Leu Asp Glu Thr Phe Ser Ser Glu Lys Ala Arg Leu
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Ala Gln Ile Thr Asn Lys Cys Thr Glu Glu Asp Leu Glu Phe Tyr Val
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                        375
    370
Arg Lys Cys Gly Asp Ile Leu Gly Val Thr Ser Lys Leu Pro Lys Asp
                    390
Gln Gln Asp Ala Lys His Ile Leu Glu His Val Phe Phe Gln Val Val
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Glu Phe Lys Lys Leu Asn Gln Glu His Asp Ile Asp Thr Ser Glu Thr
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Ala Phe Gln Asn Asn Phe
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<213> Homo sapiens
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Thr Val Tyr Ile Thr Gly Arg His Leu Asp Thr Leu Arg Val Val Ala
                            40
Gln Glu Ala Gln Ser Leu Gly Gly Gln Cys Val Pro Val Val Cys Asp
                        55
Ser Ser Gln Glu Ser Glu Val Arg Ser Leu Phe Glu Gln Val Asp Arg
                    70
Glu Gln Gln Gly Arg Leu Asp Val Leu Val Asn Asn Ala Tyr Ala Gly
                85
                                    90
Val Gln Thr Ile Leu Asn Thr Arg Asn Lys Ala Phe Trp Glu Thr Pro
                                105
                                                    110
Ala Ser Met Trp Asp Asp Ile Asn Asn Val Gly Leu Arg Gly His Tyr
                                                125
                            120
Phe Cys Ser Val Tyr Gly Ala Arg Leu Met Val Pro Ala Gly Gln Gly
                        135
Leu Ile Val Val Ile Ser Ser Pro Gly Ser Leu Gln Tyr Met Phe Asn
                                         155
                     150
Val Pro Tyr Gly Val Gly Lys Ala Ala Cys Asp Lys Leu Ala Ala Asp
                                    170
                165
Cys Ala His Glu Leu Arg Arg His Gly Val Ser Cys Val Ser Leu Trp
            180
                                185
Pro Gly Ile Val Gln Thr Glu Leu Leu Lys Glu His Met Ala Lys Glu
                            200
Glu Val Leu Gln Asp Pro Val Leu Lys Gln Phe Lys Ser Ala Phe Ser
                                             220
                         215
Ser Ala Glu Thr Thr Glu Leu Ser Gly Lys Cys Val Val Ala Leu Ala
                                         235
                     230
Thr Asp Pro Asn Ile Leu Ser Leu Ser Gly Lys Val Leu Pro Ser Cys
                                     250
                 245
Asp Leu Ala Arg Arg Tyr Gly Leu Arg Asp Val Asp Gly Arg Pro Val
                                 265
 Gln Asp Tyr Leu Ser Leu Ser Ser Val Leu Ser His Val Ser Gly Leu
                             280
        275
 Gly Trp Leu Ala Ser Tyr Leu Pro Ser Phe Leu Arg Val Pro Lys Trp
                                             300
                        295
 Ile Ile Ala Leu Tyr Thr Ser Lys Phe
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 305
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<211> 999
<212> DNA
<213> Homo sapiens
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<210> 4810
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<212> PRT
<213> Homo sapiens
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Gly Lys Ser Pro Gln Ala Asn Pro Phe Cys Glu Gln Phe Pro Ser Ala
 Val Ser Lys Ser Cys Leu Asp Ser Asp Pro Ala Gly Pro Phe Gln Gly
 Ser Gln Pro Gly Cys His Ser Gly Leu Leu Thr Asn Thr Pro Ala Ala
                             40
 Leu Val Pro Ala His Ala Arg Gln Arg Ser Gln Pro Ser Leu Leu Leu
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50
Ser Ser Ser Pro Arg Lys Ser Arg Ser Trp Gln Gly Ser Gly Pro Met
                    70
Trp Pro Gly Pro Gly Tyr Phe Pro Asp Leu Thr Ser Pro Thr Ala Gln
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Pro Leu Gln Leu Leu Gly Ala Leu His Gly Cys Ser Phe Pro Pro Pro
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            100
Leu Pro Ser Gly Gln Pro Cys Pro
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                            120
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<212> DNA
<213> Homo sapiens
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 1140
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Lys Val Thr Leu Pro Asn Tyr Asp Asn Val Pro Gly Asn Leu Met Leu
 Ser Ala Leu Gly Leu Arg Leu Gly Asp Arg Val Leu Leu Asp Gly Gln
                         55
 Lys Thr Gly Thr Leu Arg Phe Cys Gly Thr Thr Glu Phe Ala Ser Gly
                                         75
 Ser Trp Val Gly Val Glu Leu Asp Glu Pro Glu Gly Lys Asn Asp Gly
                                     90
 Ser Val Gly Gly Val Arg Tyr Phe Ile Cys Pro Pro Lys Gln Gly Leu
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                                                     110
             100
 Phe Ala Ser Val Ser Lys Ile Ser Lys Ala Val Asp Ala Pro Pro Ser
                             120
 Ser Val Thr Ser Thr Pro Gly Pro Pro Arg Met Asp Phe Ser Arg Val
                                             140
                         135
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 Thr Gly Lys Gly Arg Arg Glu His Lys Gly Lys Lys Thr Pro Ser
                                         155
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 Ser Pro Ser Leu Gly Ser Leu Gln Gln Arg Asp Gly Ala Lys Ala Glu
                                     170
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 Val Gly Asp Gln Val Leu Val Ala Gly Gln Lys Gln Gly Ile Val Arg
                                 185
             180
 Phe Tyr Gly Lys Thr Asp Phe Ala Pro Gly Tyr Trp Tyr Gly Ile Glu
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 Leu Asp Gln Pro Thr Gly Lys His Asp Gly Ser Val Phe Gly Val Arg
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 Tyr Phe Thr Cys Pro Pro Arg His Gly Val Phe Ala Pro Ala Ser Arg
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                     230
 Ile Gln Arg Ile Gly Gly Ser Thr Asp Ser Pro Gly Asp Ser Val Gly
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250
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Ala Lys Lys Val His Gln Val Thr Met Thr Gln Pro Lys Arg Thr Phe
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Thr Thr Val Arg Thr Pro Lys Asp Ile Ala Ser Glu Asn Ser Ile Ser
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Gln Ser
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Phe Gln Glu Gly Cys Leu Glu Val Gln Trp Gly Gly Arg Gly Phe Gly
                                25
Ser Pro Trp Lys Phe Leu Arg Glu Cys Ser Asn Leu Cys Leu Thr Ile
                            40
        35
Met Met Val Val Ser Trp Thr Ala Gly Gly Lys Ala Lys Pro Cys Gly
Arg Gly Gly Gly Leu Gln Arg Lys Ala Ala Ala Thr Thr Ala Ser Phe
Pro Thr His Ser His Trp Gln Thr Gly Gly Gln Val Gln Ser Pro Lys
Glu Thr Ala Ala Cys Ala Gly His Pro Pro Gly Thr Ala Phe Ser Leu
                                 105
Ile Leu Pro Val Pro Pro Thr Cys Trp Val Ser Val Ala
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<211> 528
<212> DNA
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aqcatgtcta caagctctgt acgcaaacga tctgaaggtg aagagaagac attaacaggg
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aatgetttge ccataactaa geetacatea cetgeeccag cageacagte aacaaatgge
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gttgtgaage agaagetace aggegtetat gtgeageeat ettategete tgeattaatg
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Glu Gly Glu Glu Lys Thr Leu Thr Gly Asp Val Lys Thr Ser Pro Pro
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                                 25
Arg Thr Ala Pro Lys Lys Gln Leu Pro Ser Ile Pro Lys Asn Ala Leu
Pro Ile Thr Lys Pro Thr Ser Pro Ala Pro Ala Ala Gln Ser Thr Asn
                                             60
                         55
Gly Thr His Ala Ser Tyr Gly Pro Phe Tyr Leu Glu Tyr Ser Leu Leu
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                     70
Ala Glu Phe Thr Leu Val Val Lys Gln Lys Leu Pro Gly Val Tyr Val
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                 85
 Gln Pro Ser Tyr Arg Ser Ala Leu Met
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 <212> DNA
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360
caagttaaag gcaagatcga caccatgaag aaatttaaaa gcctgttgat tcaagaactt
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1106
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Glu Asn Ile Arq Gln Leu Gly Ile Ile Val Ser Asp Phe Gln Pro Ser
            2.0
                               25
Ser Gln Ala Gly Leu Asn Gln Lys Leu Asn Phe Ile Val Thr Gly Leu
Gln Asp Ile Asp Lys Cys Arg Gln Gln Leu His Asp Ile Thr Val Pro
Leu Glu Val Phe Glu Tyr Ile Asp Gln Gly Arg Asn Pro Gln Leu Tyr
Thr Lys Glu Cys Leu Glu Arg Ala Leu Ala Lys Asn Glu Gln Val Lys
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Gly Lys Ile Asp Thr Met Lys Lys Phe Lys Ser Leu Leu Ile Gln Glu
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Leu Ser Lys Val Phe Pro Glu Asp Met Ala Lys Tyr Arg Ser Ile Arg
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Gly Glu Asp His Pro Pro Ser
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gatgtggage etgagtettg gagagaagee tteaageage attacettge atecaagaea
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ageccaaaga cetgtgacat tgttategag ggcagecaga gecctaccag eccagettet
1020
agctccccaa agccaggctc caaggctggc tcacaggagg cagaggtggg tagtgatggt
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1200
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1320
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Tyr Leu His Leu Pro Asp Leu Gly Arg Cys Ser Leu Val Cys Arg Ala
Trp Tyr Glu Leu Ile Leu Ser Leu Asp Ser Thr Arg Trp Arg Gln Leu
                        55
                                            60
Cys Leu Gly Cys Thr Glu Cys Arg His Pro Asn Trp Pro Asn Gln Pro
Asp Val Glu Pro Glu Ser Trp Arg Glu Ala Phe Lys Gln His Tyr Leu
                                    90
Ala Ser Lys Thr Trp Thr Lys Asn Ala Leu Asp Leu Glu Ser Ser Ile
                                105
Cys Phe Ser Leu Phe Arg Arg Arg Glu Arg Arg Thr Leu Ser Val
                            120
Gly Pro Gly Arg Glu Phe Asp Ser Leu Gly Ser Ala Leu Ala Met Ala
    130
                        135
Ser Leu Tyr Asp Arg Ile Val Leu Phe Pro Gly Val Tyr Glu Glu Gln
                    150
145
Gly Glu Ile Ile Leu Lys Val Pro Val Glu Ile Val Gly Gln Gly Lys
                165
                                    170
Leu Gly Glu Val Ala Leu Leu Ala Ser Ile Asp Gln His Cys Ser Thr
                                 185
            180
Thr Arg Leu Cys Asn Leu Val Phe Thr Pro Ala Trp Phe Ser Pro Ile
                            200
                                                 205
Met Tyr Lys Thr Thr Ser Gly His Val Gln Phe Asp Asn Cys Asn Phe
                        215
Glu Asn Gly His Ile Gln Val His Gly Pro Gly Thr Cys Gln Val Lys
                    230
                                         235
Phe Cys Thr Phe Lys Asn Thr His Ile Phe Leu His Asn Val Pro Leu
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255
                245
Cys Val Leu Glu Asn Cys Glu Phe Val Gly Ser Glu Asn Asn Ser Val
                                265
           260
Thr Val Glu Gly His Pro Ser Ala Asp Lys Asn Trp Ala Tyr Lys Tyr
                            280
Leu Leu Gly Leu Ile Lys Ser Ser Pro Thr Phe Leu Pro Thr Glu Asp
                        295
Ser Asp Phe Leu Met Ser Leu Asp Leu Glu Ser Arg Asp Gln Ala Trp
                                        315
                    310
Ser Pro Lys Thr Cys Asp Ile Val Ile Glu Gly Ser Gln Ser Pro Thr
                                                         335
                325
                                    330
Ser Pro Ala Ser Ser Ser Pro Lys Pro Gly Ser Lys Ala Gly Ser Gln
                                345
Glu Ala Glu Val Gly Ser Asp Gly Glu Arg Val Ala Gln Thr Pro Asp
                            360
Ser Ser Asp Gly Gly Leu Ser Pro Ser Gly Glu Asp Glu Asp Glu Asp
                        375
Gln Leu Met Tyr Arg Leu Ser Tyr Gln Val Gln Gly Pro Arg Pro Val
                                        395
Leu Gly Gly Ser Phe Leu Gly Pro Pro Leu Pro Gly Ala Ser Ile Gln
                                    410
Leu Pro Ser Cys Leu Val Leu Asn Ser Leu Gln Gln Glu Leu Gln Lys
            420
                                425
Asp Lys Glu Ala Met Ala Leu Ala Asn Ser Val Gln Gly Cys Leu Ile
                            440
Arg Lys Cys Leu Phe Arg Asp Gly Lys Gly Gly Val Phe Val Cys Ser
                        455
                                             460
    450
His Gly Arg Ala Lys Met Glu Gly Asn Ile Phe Arg Asn Leu Thr Tyr
                    470
                                        475
465
Ala Val Arg Cys Ile His Asn Ser Lys Ile Ile Met Leu Arg Asn Asp
                                     490
Ile Tyr Arg Cys Arg Ala Ser Gly Ile Phe Leu Arg Leu Glu Gly Gly
            500
                                 505
Gly Leu Ile Ala Gly Asn Asn Ile Tyr His Asn Ala Glu Ala Gly Val
                             520
                                                 525
Asp Ile Arg Lys Lys Ser Asn Pro Leu Gln Ile Gly Asn Pro Arg Ala
                                             540
Glu Phe Leu Ala Ser Arg Ala
                    550
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 gactgccage tggcgctgca gcaaaaaagg gagacgctgt ccccgctgtg cctcatcccc
 240
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aageteetge acaacegeag taacaacaag tacteetaca ecageactte agatgacaac
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<212> PRT
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Gly Arg Val Glu Val Leu Thr Asp Ala Gly Gly Trp Val Leu Ile Asp
                                     10
Arg Ser Gly Arg His Phe Gly Thr Ile Leu Asn Tyr Leu Arg Asp Gly
                                 25
Ser Val Pro Leu Pro Glu Ser Thr Arg Glu Leu Gly Glu Leu Leu Gly
                            40
Glu Ala Arg Tyr Tyr Leu Val Gln Gly Leu Ile Glu Asp Cys Gln Leu
                         55
Ala Leu Gln Gln Lys Arg Glu Thr Leu Ser Pro Leu Cys Leu Ile Pro
                                         75
65
Met Val Thr Ser Pro Arg Glu Glu Gln Gln Leu Leu Ala Ser Thr Ser
                                     90
Lys Pro Val Val Lys Leu Leu His Asn Arg Ser Asn Asn Lys Tyr Ser
                                 105
            100
Tyr Thr Ser Thr Ser Asp Asp Asn Leu Leu Lys Asn Ile Glu Leu Phe
                             120
Asp Lys Leu Ala Leu Arg Phe His Gly Arg Leu Leu Phe Leu Lys Asp
                         135
Val Leu Gly Asp Glu Ile Cys Cys Trp Ser Phe Tyr Gly Gln Gly Arg
                                         155
                     150
Lys Ile Ala Glu Val Cys Cys Thr Ser Ile Val Tyr Ala Thr Glu Lys
                                     170
Lys Gln Thr Lys Val Arg Gly Ala Pro Glu Pro Met Leu Gly Ala Gly
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                                 185
 Gly Gly His
        195
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<212> PRT
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Lys Ser Thr Gly Ser Lys Lys Ala Asn Arg Phe His Pro Tyr Ser Lys
                                 25
 Asp Lys Asn Ser Gly Thr Gly Glu Lys Lys Gly Pro Asn Arg Asn Arg
                             40
                                                 45
Val Phe Ile Ser Asn Ile Pro Tyr Asp Met Lys Trp Gln Ala Ile Lys
                         55
 Asp Leu Met Arg Glu Lys Val Gly Glu Val Thr Tyr Val Glu Leu Phe
                                         75
 Lys Asp Ala Glu Gly Lys Ser Arg Gly Cys Gly Val Val Glu Phe Lys
                85
                                     90
 Asp Glu Glu Phe Val Lys Lys Ala Leu Glu Thr Met Asn Lys Tyr Asp
                                 105
 Leu Ser Gly Arg Pro Leu Asn Ile Lys Glu Asp Pro Asp Gly Glu Asn
                             120
                                                 125
 Ala Arg Arg Ala Leu Gln Arg Thr Gly Gly Ser Phe Pro Gly Gly His
                         135
                                             140
 Val Pro Asp Met Gly Ser Gly Leu Met Asn Leu Pro Pro Ser Ile Leu
 145
                                         155
 Asn Asn Pro Asn Ile Pro Pro Glu Val Ile Ser Asn Leu Gln Ala Gly
                                     170
 Arg Leu Gly Ser Thr Ile Phe Val Ala Asn Leu Asp Phe Lys Val Gly
                                 185
 Trp Lys Lys Leu Lys Glu Val Phe Ser Ile Ala Gly Thr Val Lys Arg
                             200
                                                 205
 Ala Asp Ile Lys Glu Asp Lys Asp Gly Lys Ser Arg Gly Met Gly Thr
                         215
                                             220
 Val Thr Phe Glu Gln Ala Ile Glu Ala Val Gln Ala Ile Ser Met Phe
                     230
                                         235
 Asn Gly Gln Phe Leu Phe Asp Arg Pro Met His Val Lys Met Asp Asp
                                     250
                 245
 Lys Ser Val Pro His Glu Glu Tyr Arg Ser Pro Asp Gly Lys Thr Pro
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 Gln Leu Pro Arg Gly Leu Gly Gly Ile Gly Met Gly Leu Gly Pro Gly
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280
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Gly Gln Pro Ile Ser Ala Ser Gln Leu Asn Ile Gly Gly Val Met Gly
                                             300
                        295
Asn Leu Gly Pro Gly Gly Met Gly Met Asp Gly Pro Gly Phe Gly Gly
                                         315
                    310
Met Asn Arg Ile Gly Gly Gly Ile Gly Phe Gly Gly Leu Glu Ala Met
                                    330
                325
Asn Ser Met Gly Gly Phe Gly Gly Val Gly Arg Met Gly Glu Leu Tyr
            340
                                345
Arg Gly Ala Met Thr Ser Ser Met Glu Arg Asp Phe Gly Arg Gly Asp
                            360
        355
Ile Gly Ile Asn Arg Ala Phe Gly Asp Ser Phe Gly Arg Leu Gly Ser
                        375
Ala Met Ile Gly Gly Ile Thr Gly Arg Ile Gly Ser Ser Asn Met Gly
                                         395
                    390
Pro Val Gly Ser Gly Ile Ser Gly Gly Met Gly Ser Met Asn Ser Val
                                    410
                405
Thr Gly Gly Met Gly Met Gly Leu Asp Arg Met Ser Ser Ser Phe Asp
                                 425
Arg Met Gly Pro Gly Ile Gly Ala Ile Leu Glu Arg Ser Ile Asp Met
                            440
Asp Arq Gly Phe Leu Ser Gly Pro Met Gly Ser Gly Met Arg Glu Arg
                        455
                                             460
Ile Gly Ser Lys Gly Asn Gln Ile Phe Val Arg Asn Leu Pro Phe Asp
                    470
                                         475
Leu Thr Trp Gln Lys Leu Lys Glu Lys Phe Ser Gln Cys Gly His Val
                485
                                     490
Met Phe Ala Glu Ile Lys Met Glu Asn Gly Lys Ser Lys Gly Cys Gly
                                 505
Thr Val Arg Phe Asp Ser Pro Glu Ser Ala Glu Lys Ala Cys Arg Ile
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Met Asn Gly Ile Lys Ile Ser Gly Arg Glu Ile Asp Val Arg Leu Asp
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Arg Asn Ala
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<210> 4830

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Val Gly Asp Met Val Leu Leu Glu Pro Leu Asn Glu Glu Thr Phe Ile
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Asn Asn Leu Lys Lys Arg Phe Asp His Ser Glu Ile Tyr Thr Tyr Ile
                            40
Gly Ser Val Val Ile Ser Val Asn Pro Tyr Arg Ser Leu Pro Ile Tyr
                        55
Ser Pro Glu Lys Val Glu Glu Tyr Arg Asn Arg Asn Phe Tyr Glu Leu
                    70
                                        75
Ser Pro His Ile Phe Ala Leu Ser Asp Glu Ala Tyr Arg Ser Leu Arg
                                    90
Asp Gln Asp Lys Asp Gln Cys Ile Leu Ile Thr Gly Glu Ser Gly Ala
                                105
Gly Lys Thr Glu Ala Ser Lys Leu Val Met Ser Tyr Val Ala Ala Val
                           120
Cys Gly Lys Gly Ala Glu Val Asn Gln Val Lys Glu Gln Leu Leu Gln
                        135
Ser Asn Pro Val Leu Glu Ala Phe Gly Asn Ala Lys Thr Val Arg Asn
                                        155
                    150
Asp Asn Ser Ser Arg Phe Gly Lys Tyr Met Asp Ile Glu Phe Asp Phe
                                    170
Lys Gly Asp Pro Leu Gly Gly Val Ile Ser Asn Tyr Leu Leu Glu Lys
                                185
Ser Arg Val Val Lys Gln Pro Arg Gly Glu Arg Asn Phe His Val Phe
                            200
Tyr Gln Leu Leu Ser Gly Ala Ser Glu Glu Leu Leu Asn Lys Leu Lys
                                            220
                        215
Leu Glu Arg Asp Phe Ser Arg Tyr Asn Tyr Leu Ser Leu Asp Ser Ala
                                        235
                   230
Lys Val Asn Gly Val Asp Asp Ala Ala Asn Phe Arg Thr Val Arg Asn
                                    250
Ala Met Gln Ile Val Gly Phe Met Asp His Glu Ala Glu Ser Val Leu
                                265
Ala Val Val Ala Ala Val Leu Lys Leu Gly Asn Ile Glu Phe Lys Pro
                            280
Glu Ser Arg Val Asn Gly Leu Asp Glu Ser Lys Ile Lys Asp Lys Asn
                        295
Glu Leu Lys Glu Ile Cys Glu Leu Thr Gly Ile Asp Gln Ser Val Leu
                                        315
                    310
Glu Arg Ala Phe Ser Phe Arg Thr Val Glu Ala Lys Gln Glu Lys Val
                                    330
Ser Thr Thr Leu Asn Val Ala Gln Ala Tyr Tyr Ala Arg Asp Ala Leu
                                345
Ala Lys Asn Leu Tyr Ser Arg Leu Phe Ser Trp Leu Val Asn Arg Ile
                            360
Asn Glu Ser Ile Lys Ala Gln Thr Lys Val Arg Lys Lys Val Met Gly
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380
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                       375
Val Leu Asp Ile Tyr Gly Phe Glu Ile Phe Glu Asp Asn Ser Phe Glu
                    390
                                       395
Gln Phe Ile Ile Asn Tyr Cys Asn Glu Lys Leu Gln Gln Ile Phe Ile
                                   410
                405
Glu Leu Thr Leu Lys Glu Glu Glu Glu Glu Tyr Ile Arg Glu Asp Ile
                               425
Glu Trp Thr His Ile Asp Tyr Phe Asn Asn Ala Ile Ile Cys Asp Leu
                           440
Ile Glu Asn Asn Thr Asn Gly Ile Leu Ala Met Leu Asp Glu Glu Cys
                        455
Leu Arg Pro Gly Thr Val Thr Asp Glu Thr Phe Leu Glu Lys Leu Asn
                    470
                                        475
465
Gln Val Cys Ala Thr His Gln His Phe Glu Ser Arg Met Ser Lys Cys
                                    490
Ser Arg Phe Leu Asn Asp Thr Ser Leu Pro His Ser Cys Phe Arg Ile
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<211> 578
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<213> Homo sapiens
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cagcagageg tgeteggegg ccaggaccag etgegegtee gtgtgaegga getggaggae
gaggtgcgca acctgcgcaa gatcaatcgg gacctgttcg acttctccac gcgcttcatc
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teagttetgt gtegtgtteg ggttttteet etgtgaetgg geegtettgg tgtetegtgg
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Ala Pro Val Ser Met Leu Ser Ser Asp Phe Arg Pro Ser Leu Pro Leu
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25
            20
Pro His Phe Asn Lys His Leu Leu Gly Ala Glu His Gly Asp Glu Pro
Arg His Gly Gly Leu Thr Leu Arg Leu Gly Leu His Gln Gln Ser Val
Leu Gly Gly Gln Asp Gln Leu Arg Val Arg Val Thr Glu Leu Glu Asp
Glu Val Arg Asn Leu Arg Lys Ile Asn Arg Asp Leu Phe Asp Phe Ser
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                                                         95
Thr Arg Phe Ile Thr Arg Pro Ala Lys
            100
                                105
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<2115 872
<212> DNA
<213> Homo sapiens
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ctcaacaact gagatgaacg tcgactcgct tgcaggcaag ttgtcactca gcagcgatct
qaactatatc ctgggttcca gaaaaggcag aggttcttac cgaaagcagg ggaggaagcc
quaquecaag gaggtegtea ettgeeggga aggtggeteg ggeeaggetg cacteaaaac
ccgtgctctg tccacactgc tacggggcca gagccaagga agcttccact tcttccccca
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atqcagcgga acctgaagat ggagatgett gccgggtgtg gggctgggat gtgccaggtc
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cqtccatcat cagggctcgg cctcagcacc ct
872
<210> 4834
<211> 147
<212> PRT
<213> Homo sapiens
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Val Ala Gly Leu Val Gly Val Thr Cys Val Phe Pro Ile Asp Leu Ala
Lys Thr Arq Leu Gln Asn Gln His Gly Lys Ala Met Tyr Lys Gly Met
Ile Asp Cys Leu Met Lys Thr Ala Arg Ala Glu Gly Phe Phe Gly Met
Tyr Arg Gly Ala Ala Val Asn Leu Thr Leu Val Thr Pro Glu Lys Ala
65
                                        75
Ile Lys Leu Ala Ala Asn Asp Phe Phe Arg Arg Leu Leu Met Glu Asp
Gly Met Gln Arg Asn Leu Lys Met Glu Met Leu Ala Gly Cys Gly Ala
Gly Met Cys Gln Val Val Thr Cys Pro Met Glu Met Leu Lys Ile
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Gln Leu Gln Ala Cys Trp Thr Pro Gly Arg Pro Ser Ser Gly Leu Gly
                        135
                                            140
Leu Ser Thr
145
<210> 4835
<211> 1846
<212> DNA
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ctgcacaaag ctttcgcccg agctgaactg gaacgcacgt accaggagat ccaggagtta
cagtgggaga tccagaatac cagccatctg gccgttgatg gggaccgggc agctgcttgg
cccgtgggta ttccagcacc atcccgcccg gcctcccgct ttgaggtgct gcgctgggac
tacttcacgg agcagcacgc tttctcctgc gccgatggct caccccgctg cccactgcgt
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cgctaccacc cggccttgcg gctccagaag cagcagctgg tgaatggcta ccgacgcttt
gateeggeec qqqqtatqqa atacacqetq qaettqcaqe tqqaqqcaet gaeeceecag
ggaggccgcc ggcccctcac tcgccgagtg cagctgctcc ggccgctgag ccgcgtggag
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cetggtgatg etgeggeage cetgaceetg etgetaetgt atgageegeg eeaggeeeag
cgcgtggccc atgcagatgt cttcgcacct gtcaaggccc acgtggcaga gctggagcgg
780
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90
                85
Cys Pro Leu Arg Gly Ala Asp Arg Ala Asp Val Ala Asp Val Leu Gly
                                105
            100
Thr Ala Leu Glu Glu Leu Asn Arg Arg Tyr His Pro Ala Leu Arg Leu
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Gln Lys Gln Gln Leu Val Asn Gly Tyr Arg Arg Phe Asp Pro Ala Arg
                        135
Gly Met Glu Tyr Thr Leu Asp Leu Gln Leu Glu Ala Leu Thr Pro Gln
                    150
                                        155
Gly Gly Arg Arg Pro Leu Thr Arg Arg Val Gln Leu Leu Arg Pro Leu
                165
                                    170
Ser Arg Val Glu Ile Leu Pro Val Pro Tyr Val Thr Glu Ala Ser Arg
                                185
            180
Leu Thr Val Leu Leu Pro Leu Ala Ala Ala Glu Arg Asp Leu Ala Pro
                            200
Gly Phe Leu Glu Ala Phe Ala Thr Ala Ala Leu Glu Pro Gly Asp Ala
                        215
                                            220
    210
Ala Ala Ala Leu Thr Leu Leu Leu Tyr Glu Pro Arg Gln Ala Gln
                    230
                                        235
Arg Val Ala His Ala Asp Val Phe Ala Pro Val Lys Ala His Val Ala
                245
                                    250
Glu Leu Glu Arg Arg Phe Pro Gly Ala Arg Val Pro Trp Leu Ser Val
                                                    270
                                265
Gln Thr Ala Ala Pro Ser Pro Leu Arg Leu Met Asp Leu Leu Ser Lys
        275
                            280
                                                285
Lys His Pro Leu Asp Thr Leu Phe Leu Leu Ala Gly Pro Asp Thr Val
                        295
                                            300
Leu Thr Pro Asp Phe Leu Asn Arg Cys Arg Met His Ala Ile Ser Gly
                                        315
                                                            320
                    310
Trp Gln Ala Phe Phe Pro Met His Phe Gln Ala Phe His Pro Ala Val
                325
                                    330
Ala Pro Pro Gln Gly Pro Gly Pro Pro Glu Leu Gly Pro
            340
                                345
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actgtaaatt atgatagtgt caattctgac aactctaagc caaagatatt taaaagtcaa
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attcaccett qtqqaaatce tacaqtqatt qaqqacqctt tqqacaaqat taaaagcaat
gaccetgaca ccacagaagt caatttgaac aacattgaga acatcacaac acagaccett
accegetttg etgaageest caaggacaac actgtggtga agaegtteag tetggeeaac
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acgcatgccg acgacagtgc agccatggcc attgcagaga tgctcaaagt caatgagcac
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Gln Glu Glu Glu Glu Glu Glu Asp Ser Asp Glu Gly Glu Arg Thr Ile
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Glu Thr Ala Lys Gly Ile Asn Gly Thr Val Asn Tyr Asp Ser Val Asn
                            40
Ser Asp Asn Ser Lys Pro Lys Ile Phe Lys Ser Gln Ile Glu Asn Ile
                        55
Asn Leu Thr Asn Gly Ser Asn Gly Arg Asn Thr Glu Ser Pro Ala Ala
                    70
                                        75
Ile His Pro Cys Gly Asn Pro Thr Val Ile Glu Asp Ala Leu Asp Lys
                                    90
Ile Lys Ser Asn Asp Pro Asp Thr Thr Glu Val Asn Leu Asn Asn Ile
                                105
            100
Glu Asn Ile Thr Thr Gln Thr Leu Thr Arg Phe Ala Glu Ala Leu Lys
        115
                            120
                                                125
Asp Asn Thr Val Val Lys Thr Phe Ser Leu Ala Asn Thr His Ala Asp
                        135
                                            140
Asp Ser Ala Ala Met Ala Ile Ala Glu Met Leu Lys Val Asn Glu His
                                        155
                    150
Ile Thr Asn Val Asn Val Glu Ser Asn Phe Ile Thr Gly Lys Gly Ile
                                    170
Leu Ala Ile Met Arg Ala Leu Gln His Asn Thr Val Leu Thr Glu Leu
                                185
                                                     190
Arg Phe His Asn Gln Arg His Ile Met Gly Ser Gln Val Glu Met Glu
Ile Val Lys Leu Leu Lys Glu Asn Thr Thr Leu Leu Arg Leu Gly Tyr
                        215
His Phe Glu Leu Pro Gly Pro Arg Met Ser Met Thr Ser Ile Leu Thr
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225
                    230
                                        235
Arg Asn Met Asp Lys Gln Arg Gln Lys Arg Leu Gln Glu Gln Lys Gln
                                    250
                245
Gln Glu Gly Tyr Asp Gly Gly Pro Asn Leu Arg Thr Lys Val Trp Gln
                                265
Arg Gly Thr Pro Ser Pro Ser Pro Tyr Val Ser Pro Arg His Ser Pro
                            280
Trp Ser Ser Pro Lys Leu Pro Tyr Gly Glu Thr Thr Arg
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1140
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Gly Thr Pro Ala Arg Gln Lys Leu Glu Lys Ala Arg Asp Val Ala Arg
Asp Pro Gly Thr Ser Pro Ser Ser Pro Gly Pro Pro Gly Pro Asp
       35
Gly His Ser Arg Tyr Ser Ala His Ser Val Leu Gly His Pro Ala Pro
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Ala Val
65
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<212> DNA
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qaggageget caaggecage gaggaaagge teageageaa gttgaetgea etagagegge
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<213> Homo sapiens

<210> 4843

<211> 6403

<212> DNA

<213> Homo sapiens

<400> 4843

840

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Asn Lys Asp Thr Gly His Ser Asn Arg Gln Ser Asp Val Arg Ile Lys
Phe Glu His Asn Gly Glu Arg Arg Ile Ile Ala Phe Ser Arg Pro Val
Lys Tyr Glu Asp Val Glu His Lys Val Thr Thr Val Phe Gly Gln Pro
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Gln Asp Asp Leu Asp Lys Ala Ile Asp Ile Leu Asp Arg Ser Ser Ser
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Arg Ser Arg His Leu Ser Val Ser Ser Gln Asn Pro Gly Arg Ser Ser
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Pro Pro Pro Gly Tyr Val Pro Glu Arg Gln Gln His Ile Ala Arg Gln
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Gly Thr Tyr Pro Arg Arg Tyr His Val Ser Val His His Lys Asp Tyr
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Ser Asp Gly Arg Arg Thr Phe Pro Arg Ile Arg Arg His Gln Gly Asn
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Ser Ala Asp Ser Glu Asn Ala Leu Ser Val Gln Glu Arg Asn Val Pro
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Thr Lys Ser Pro Ser Ala Pro Ile Asn Trp Arg Arg Gly Lys Leu Leu
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Gly Gln Gly Ala Phe Gly Arg Val Tyr Leu Cys Tyr Asp Val Asp Thr
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Gly Arg Glu Leu Ala Ser Lys Gln Val Gln Phe Asp Pro Asp Ser Pro
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Lvs Asn Leu Gln His Glu Arg Ile Val Gln Tyr Tyr Gly Cys Leu Arg
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Gly Ser Val Lys Asp Gln Leu Lys Ala Tyr Gly Ala Leu Thr Glu Ser
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Val Thr Arg Lys Tyr Thr Arg Gln Ile Leu Glu Gly Met Ser Tyr Leu
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Arg Asp Ser Ala Gly Asn Val Lys Leu Gly Asp Phe Gly Ala Ser Lys
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Ser Glu His Gly Arg Asp Phe Leu Arg Arg Ile Phe Val Glu Ala Arg
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Leu Arg Met Leu Pro Ile Ser Gly Thr Cys Lys Ala Phe Leu Glu Asp
Met Lys Lys Tyr Ala Glu Thr Phe Leu Glu Pro Trp Phe Lys Ala Pro
Asn Lys Gly Thr Phe Gln Ile Val Tyr Lys Ser Arg Asn Asn Ser His
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Val Asn Arg Glu Glu Val Ile Arg Glu Leu Ala Gly Ile Val Cys Thr
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Leu Asn Ser Glu Asn Lys Val Asp Leu Thr Asn Pro Gln Tyr Thr Val
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Val Val Glu Ile Ile Lys Ala Val Cys Cys Leu Ser Val Val Lys Asp
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Tyr Met Leu Phe Arg Lys Tyr Asn Leu Gln Glu Val Val Lys Ser Pro
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Lys Asp Pro Ser Gln Leu Asn Ser Lys Gln Gly Asn Gly Lys Glu Ala
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Lys Leu Glu Ser Ala Asp Lys Ser Asp Gln Asn Asn Thr Ala Glu Gly
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Lys Asn Asn Gln Gln Val Pro Glu Asn Thr Glu Glu Leu Gly Gln Thr
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Lys Pro Thr Ser Asn Pro Gln Val Val Asn Glu Gly Gly Ala Lys Pro
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Gln Glu Arg Gly Ser Ala His Leu Val Ala Leu Lys Cys Ile Pro Lys
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Lys Ala Leu Arg Gly Lys Glu Ala Leu Val Glu Asn Glu Ile Ala Val
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Ala Leu Pro Asp Gln Tyr Gln Glu Asp Ala Ser Asp Met Lys Asp Met
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Ser Lys Tyr Lys Pro His Ile Leu Leu Ser Gln Glu Asn Thr Gln Ile
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Arg Asp Leu Gln Gln Glu Asn Arg Glu Leu Trp Ile Ser Leu Glu Glu
His Gln Asp Ala Leu Glu Leu Ile Met Ser Lys Tyr Arg Lys Gln Met
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Leu Gln Leu Met Val Ala Lys Lys Ala Val Asp Ala Glu Pro Val Leu
                            120
Lvs Ala His Gln Ser His Ser Ala Glu Ile Glu Ser Gln Ile Asp Arq
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Ile Cys Glu Met Gly Glu Val Met Arg Lys Ala Val Gln Val Asp Asp
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Asp Gln Phe Cys Lys Ile Gln Glu Lys Leu Ala Gln Leu Glu Leu Glu
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Asn Lys Glu Leu Arq Glu Leu Leu Ser Ile Ser Ser Glu Ser Leu Gln
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Ala Arg Lys Glu Asn Ser Met Asp Thr Ala Ser Gln Ala Ile Lys
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4039

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Arg Lys Val Glu Leu Pro Val Pro Thr His Arg Arg Pro Val Gln Ala
Trp Val Glu Ser Leu Arg Gly Phe Glu Gln Glu Arg Val Gly Leu Ala
Asp Leu His Pro Asp Val Phe Ala Thr Ala Pro Arg Leu Asp Ile Leu
His Gln Val Ala Met Trp Gln Lys Asn Phe Lys Arg Ile Ser Tyr Ala
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Lys Thr Lys Thr Arg Ala Glu Val Arg Gly Gly Arg Lys Pro Xaa
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Ala Leu Ala Arg Arg Cys Cys Pro Trp Pro Pro Gly Pro Thr Ser
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Tyr Tyr Met Leu Pro Met Lys Val Arg Ala Leu Gly Leu Lys Val
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Tyr Arg Arg Trp Gly Asp Ser Val Leu Leu Val Asp Leu Thr His Glu
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<400> 4856
Met Ala Phe Asn Phe Gly Ala Pro Ser Gly Thr Ser Gly Thr Ala Ala
Ala Thr Ala Ala Pro Ala Gly Gly Phe Gly Phe Gly Thr Thr Ser
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Thr Thr Ala Gly Ser Ala Phe Ser Phe Ser Ala Pro Thr Asn Thr Gly
        35
Thr Thr Gly Leu Phe Gly Gly Thr Gln Asn Lys Gly Phe Gly Phe Gly
                        55
Thr Gly Phe Gly Thr Thr Thr Gly Thr Ser Thr Gly Leu Gly Thr Gly
Leu Gly Thr Gly Leu Gly Phe Gly Gly Phe Asn Thr Gln Gln Gln
Gln Gln Thr Thr Leu Gly Gly Leu Phe Ser Gln Pro Thr Gln Ala Pro
            100
                                105
Thr Gln Ser Asn Gln Leu Ile Asn Thr Ala Ser Ala Leu Ser Ala Pro
                            120
Thr Leu Leu Gly Asp Glu Arg Asp Ala Ile Leu Ala Lys Trp Asn Gln
                        135
                                            140
Leu Gln Ala Phe Trp Gly Thr Gly Lys Gly Tyr Phe Asn Asn Asn Ile
                                        155
145
                    150
Pro Pro Val Glu Phe Thr Gln Glu Asn Pro Phe Cys Arg Phe Lys Ala
                                    170
                165
Val Gly Tyr Ser Cys Met Pro Ser Asn Lys Asp Glu Asp Gly Leu Val
            180
                                185
Val Leu Val Phe Asn Lys Lys Glu Thr Glu Ile Arg Ser Gln Gln Gln
                            200
                                                205
Gln Leu Val Glu Ser Leu His Lys Val Leu Gly Gly Asn Gln Thr Leu
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Thr Val Asn Val Glu Gly Thr Lys Thr Leu Pro Asp Asp
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235

225

230

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acaaaggttc taaaatgtga agagtttgtt tgaaaaatag tttgtagacc attttattta aatatatgaa caaccaatgg gctactgcaa tccaagtaaa ctcttcacat tttagaacct ttgtgaagta tagtaagata aagtaagact gttggtcttt ggcagattcc tcctgccccc 1560 cagacaggga catagatata cagataaatg tttatatagt taaagagcgg aggcccaggt gaaaattccc caccccaagc tggctcccca acccaaaaat taccattggc ccctgagaac 1680 acccaaaqac caccctccca qatttcacac aatatcaatt gcaggaacag tcgtgcatgg caaagtatag ttctatgcat ttcacagcac agaaaaccct tctttcagag ggcatgcaag 1800 ttqccqqaqa ttacaccccc aaqtctggtg ctcttgtgat gagcaactta ctggcaaaca catctcccaq tqttgtaagc aggcaaatac attgagcaca ttttgctgta attccatcta 1920 tttqcaatqc ctqcacagtg tctqtctctg gctgttaact tactcattct tgacagaact ctqctttatt gattgcactt ttttaaaaat gccaaaggca ttttcacact tgttagcttg 2040 acceptacca ctagggtata agagcaagtg ttetetatge ettaettget acagteteet 2100 cettetttqq aqaqqattcc cqtaatqqta acqqtqatqq gqqagqctqq tgccaacgac acaaatacat ttctgtggtg gaaaggtacg gcaaatcttc tatctcactt gagaaggctt teteettggg atgggtgetg ggteeetttt ggggagtgga gagtettggt ggqqtgteea ctgaagaccg gggcttttct ggggtttctt ggctcctcaa ttcacgttta caaacagttt 2340 cagataagga accacagggt tcctctttaa taggagagtg cttaggagtt tttgttttga 2400 cttttqaaaa ttcaggagcc agctttttaa caggagtctt tctttctgta cttccaaatg 2460 ggggtagaag totaacccct ccaccccctc toctccccag cagtcccacg cgggtatggg 2520 agagaatgaa gttctttgtc tctaagggat tcaaaccaga aacggaggga cctctggttc ccagagggag gaaaatccat gatgtetget geecagggag ctattgecac egeeteettg qqatqaaqta ttqccaqcta ccaacagttc cttcccaacg gccatcttcc agccttctta aacgactoot agcatottog ggaggotoot gaaggactga agcaaaggaa atototgaag ggatttagtc cttqaaaqqq aqtaqqqata cttaqqqtqt tctqtqttqa gcgcttcttc ctatctctcc agettcatgt atgtgtgtct ttatgtccaa gcaattgagc caacaagtcc 2880 tcagaat 2887

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<212> PRT
<213> Homo sapiens
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Gly Trp Trp Arg Leu Gly Ser Ser Ser Gln Ala Ala Cys Leu Lys Gln
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Ile Leu Leu Cln Leu Asp Leu Ile Glu Gln Gln Gln Gln Leu
                            40
Gln Ala Lys Glu Lys Glu Ile Glu Glu Leu Lys Ser Glu Arg Asp Thr
                        55
Leu Leu Ala Arg Ile Glu Arg Met Glu Arg Arg Met Gln Leu Val Lys
                                        75
Lys Asp Asn Glu Lys Glu Arg His Lys Leu Phe Gln Gly Tyr Glu Thr
Glu Glu Arg Glu Glu Thr Glu Leu Ser Glu Lys Ile Lys Leu Glu Cys
            100
                                105
Gln Pro Glu Leu Ser Glu Thr Ser Gln Thr Leu Pro Pro Lys Pro Phe
                            120
        115
Ser Cys Gly Arg Ser Gly Lys Gly His Lys Arg Lys Ser Pro Phe Gly
                        135
                                            140
Ser Thr Glu Arg Lys Thr Pro Val Lys Lys Leu Ala Pro Glu Phe Ser
                   150
                                        155
Lys Val Lys Thr Lys Thr Pro Lys His Ser Pro Ile Lys Glu Glu Pro
                165
                                    170
                                                        175
Cys Gly Ser Leu Ser Glu Thr Val Cys Lys Arg Glu Leu Arg Ser Gln
            180
                                185
Glu Thr Pro Glu Lys Pro Arg Ser Ser Val Asp Thr Pro Pro Arg Leu
                                                205
                            200
Ser Thr Pro Gln Lys Gly Pro Ser Thr His Pro Lys Glu Lys Ala Phe
                        215
Ser Ser Glu Ile Glu Asp Leu Pro Tyr Leu Ser Thr Thr Glu Met Tyr
                    230
                                        235
Leu Cys Arg Trp His Gln Pro Pro Pro Ser Pro Leu Pro Leu Arg Glu
                245
                                    250
Ser Ser Pro Lys Lys Glu Glu Thr Val Ala Ser Lys Ala
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                                265
<210> 4859
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<212> DNA
<213> Homo sapiens
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gggeteette aggagetete tacccagggg caaggagage ccagagagaa gegeeetggt

240

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ctcttgaget teetgatetg etcetgteec cegeteteet ceaetecett geettteect
aggttgtccc ctccctgggc ttttgtgtgt tttgggagat gtcacctaac caggacattg
atattcaatc ccatcccct tcctcccacc ctgccccact ttgatttaat cctttqqctg
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acaqaqacct gtccttgtgc agtctgcacc ctgcactccc tcccttgcct gtagatgttc
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<210> 4860
<211> 173
<212> PRT
<213> Homo sapiens
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Trp Thr Leu Asp Leu Glu Pro Arg Gly Pro Val His Ile His Pro Thr
Arg Val Ser Gly Gly Leu Pro Arg Cys Leu Cys Trp Val Ala Val Val
        35
Val Pro Arg Gly Met Glu Cys Pro Gly Leu Leu Gln Glu Leu Ser Thr
                        55
Gln Gly Gln Gly Glu Pro Arg Glu Lys Arg Pro Gly Leu Leu Ser Phe
Leu Ile Cys Ser Cys Pro Pro Leu Ser Ser Thr Pro Leu Pro Phe Pro
                85
Arg Leu Ser Pro Pro Trp Ala Phe Val Cys Phe Gly Arg Cys His Leu
            100
                                 105
Thr Arg Thr Leu Ile Phe Asn Pro Ile Pro Leu Pro Pro Thr Leu Pro
                            120
His Phe Asp Leu Ile Leu Trp Leu Trp Ala Glu Ala Ser Gln Gly Ser
                        135
Trp Val Gly Trp Val Leu Arg Pro Pro Gln Thr Ser Thr Glu Thr Cys
                                                             160
                    150
                                        155
145
Pro Cys Ala Val Cys Thr Leu His Ser Leu Pro Cys Leu
                                     170
                165
<210> 4861
<211> 1622
<212> DNA
<213> Homo sapiens
<400> 4861
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ctggtgtgtg tttcctttca gcgtcaaggg ttccacactg ttgggagtcg ctgcaagaat 120 cggacaggcg ctgagcacct gtggctgacc cgacatctca gggacccatt tgtgaaggct gcgaaggtgg agagttaccg gtgtcgaagc gccttcaagc tcctggaggt gaacgagagg caccagattc tgcggcccgg ccttcgggtg ttagactgtg gggcagctcc tggggcctgg agtcaggtgg cggtgcagaa ggtcaacgcc gcaggcacag atcccagctc tcctgttggc ttcgtgcttg gggtagatct tcttcacata ttccccctgg aaggagcaac ttttctgtgc cetgetgacg tgactgacce gagaacetca cagagaatee tegaggtget teetggcagg agagcagatg tgattetgag egacatggeg eccaatgeca cagggtteeg ggacetegat catgacagge teateageet gtgcctgace etteteageg tgaccccaga catcetgcaa cctgggggga cattcctttg taaaacctgg gctggaagtc aaagccgtcg gttacagagg agactgacag aggaattcca gaatgtaagg atcatcaaac ctgaagccag caggaaagag 720 tcatcagaag tgtacttctt ggccacacag taccacggaa ggaagggcac tgtgaagcag tgaggatttc ttgtgccatt ttcataatgg tcattagctc cttttaagct agaaacgtag cctgagctcc tgaagagttc ctgggagatt tgagctgatt ttggagatgg agcaggacaa qtqqqqqtc tctctctct tttctctctc tctctttta accaaaaaga gatgacaaaa ctaagttcag gggccatgga aaatgaaaaa gtccgctata ttgtgatttg ggaagagaaa gttatcaaga gaaagaggtg aggatggaag gatggagaaa aacagactgt gggaaggatc agaaggaatc cgccgaggca gggatgggtg tgcccatgtg tgccttgacg ggacttcatc ttatagactg ttaaactgtc acacacaaac aggettteca eccetgetet gagageacca cgcacagatt tccagttctt agtgtggctg tttaaaagtag aaaatctggg ggctgggtga ggccactcat gcctgtaaac ccagggcttt agaaggctga ggctgggggga ttgcttgaag tcaggagttc aagaccaacc tgggcaacat agcaacaccc cccatgtcta caaaaatgaa aaaccaaaaa gcaaaccaaa agaaaaatct gaaatttcca tctggggatt aacttctgtc tttctggtga acaatatagc aattcacgca ttcttcaagc agcaaaagtt cccggaacaa ttagggaaga cgtatggtct gaatttatcc aggcagtggg tctgctttgg tttttgctgg aaatttatat cagtgtctgg gctcccaaga acataaatgt aattgccaaa gcaaaaaaa 1620 aa 1622

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<211> 260
<212> PRT
<213> Homo sapiens
<400> 4862
Leu Gln Thr Ser Gly Gly Ala Leu Gln Ala Arg Gly Thr Pro Met Ala
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Gly Tyr Leu Lys Leu Val Cys Val Ser Phe Gln Arg Gln Gly Phe His
                                25
Thr Val Gly Ser Arg Cys Lys Asn Arg Thr Gly Ala Glu His Leu Trp
                            40
Leu Thr Arg His Leu Arg Asp Pro Phe Val Lys Ala Ala Lys Val Glu
                                            60
                        55
Ser Tyr Arg Cys Arg Ser Ala Phe Lys Leu Leu Glu Val Asn Glu Arg
                    70
                                        75
His Gln Ile Leu Arg Pro Gly Leu Arg Val Leu Asp Cys Gly Ala Ala
                                    90
Pro Gly Ala Trp Ser Gln Val Ala Val Gln Lys Val Asn Ala Ala Gly
                                105
Thr Asp Pro Ser Ser Pro Val Gly Phe Val Leu Gly Val Asp Leu Leu
                            120
                                                 125
His Ile Phe Pro Leu Glu Gly Ala Thr Phe Leu Cys Pro Ala Asp Val
                        135
                                             140
Thr Asp Pro Arg Thr Ser Gln Arg Ile Leu Glu Val Leu Pro Gly Arg
                    150
                                        155
Arg Ala Asp Val Ile Leu Ser Asp Met Ala Pro Asn Ala Thr Gly Phe
                165
                                    170
                                                        175
Arg Asp Leu Asp His Asp Arg Leu Ile Ser Leu Cys Leu Thr Leu Leu
                                                     190
                                185
Ser Val Thr Pro Asp Ile Leu Gln Pro Gly Gly Thr Phe Leu Cys Lys
                                                 205
                            200
Thr Trp Ala Gly Ser Gln Ser Arg Arg Leu Gln Arg Arg Leu Thr Glu
                        215
                                             220
Glu Phe Gln Asn Val Arg Ile Ile Lys Pro Glu Ala Ser Arg Lys Glu
                                        235
                    230
Ser Ser Glu Val Tyr Phe Leu Ala Thr .Gln Tyr His Gly Arg Lys Gly
                245
                                    250
                                                         255
Thr Val Lys Gln
            260
<210> 4863
<211> 355
<212> DNA
<213> Homo sapiens
<400> 4863
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gececaaata teacagecaa ceteaceteg tecetgetga gegtetgtgg gtggagecag
accatcaacc ctgaggacga cacggatect ggccatgctg acctggtect ctatatcact
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aggtttgacc tggagttgcc tgatggtaac neggeagtgc ggggegteac ceagetggge
ggggcctgct ccccaacctg gagctgcctc attaccgagg acactggctt cgacctggga
gtcaccattg cccatgagat tgggcacagc ttcggcctgg agcacgacgg cgcgc
355
<210> 4864
<211> 118
<212> PRT
<213> Homo sapiens
<400> 4864
Leu Gly Ala His Phe Arg Val His Leu Val Lys Met Val Ile Leu Thr
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1
Glu Pro Glu Gly Ala Pro Asn Ile Thr Ala Asn Leu Thr Ser Ser Leu
                                25
Leu Ser Val Cys Gly Trp Ser Gln Thr Ile Asn Pro Glu Asp Asp Thr
                            40
Asp Pro Gly His Ala Asp Leu Val Leu Tyr Ile Thr Arg Phe Asp Leu
                        55
                                            60
Glu Leu Pro Asp Gly Asn Xaa Ala Val Arg Gly Val Thr Gln Leu Gly
                                        75
Gly Ala Cys Ser Pro Thr Trp Ser Cys Leu Ile Thr Glu Asp Thr Gly
                                    90
Phe Asp Leu Gly Val Thr Ile Ala His Glu Ile Gly His Ser Phe Gly
            100
                                105
                                                     110
Leu Glu His Asp Gly Ala
        115
<210> 4865
<211> 444
<212> DNA
<213> Homo sapiens
<400> 4865
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aaggeetteg eegacagete ttacetgett egecaceage geacteacte tggecagaag
ccctacaagt gcccacattg tggcaaggcc tteggcgaca gctcctacct cctgcgacac
caqeqcacce acagecacga geggecetae agetgeaceg agtgeggeaa gtgetatage
300
cagaactegt ecetgegeag ccatcagagg gtgcacaceg gtcagaggec ettcagetgt
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444
<210> 4866
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<211> 148
<212> PRT
<213> Homo sapiens
<400> 4866
Thr Gly Glu Lys Pro Tyr Lys Cys Glu Val Cys Ser Lys Ala Phe Ser
Gln Ser Ser Asp Leu Ile Lys His Gln Arg Thr His Thr Gly Glu Arg
Pro Tyr Lys Cys Pro Arg Cys Gly Lys Ala Phe Ala Asp Ser Ser Tyr
Leu Leu Arg His Gln Arg Thr His Ser Gly Gln Lys Pro Tyr Lys Cys
Pro His Cys Gly Lys Ala Phe Gly Asp Ser Ser Tyr Leu Leu Arg His
                    70
                                        75
Gln Arg Thr His Ser His Glu Arg Pro Tyr Ser Cys Thr Glu Cys Gly
                                     90
Lys Cys Tyr Ser Gln Asn Ser Ser Leu Arg Ser His Gln Arg Val His
                                105
Thr Gly Gln Arg Pro Phe Ser Cys Gly Ile Cys Gly Lys Ser Phe Ser
        115
                            120
                                                125
Gln Arg Ser Ala Leu Ile Pro His Ala Arg Ser His Ala Arg Glu Lys
                        135
                                            140
    130
Pro Phe Thr Arg
145
<210> 4867
<211> 391
<212> DNA
<213> Homo sapiens
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cettetecae atececatte tggtaggaaa agteacecat gecaggatat ceccagecea
gagacagece cagggggtge tgcctggaga cagecgggat agettcagte teetgaceet
gacacgggct gcaccaccag acaatgggca ttttcaggcc agactctggc acaaagagaa
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agegetetae teccataget ecceaetgta t
391
<210> 4868
<211> 125
<212> PRT
<213> Homo sapiens
<400> 4868
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Met Gly Val Glu Arg Tyr Leu Leu His Pro Ser Gln Leu Leu Arg Ser

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Leu Trp Ala Ile Ala Leu Ala Leu Pro Leu Leu Phe Val Pro Glu Ser
Gly Leu Lys Met Pro Ile Val Trp Trp Cys Ser Pro Cys Gln Gly Gln
                            40
Glu Thr Glu Ala Ile Pro Ala Val Ser Arg Gln His Pro Leu Gly Leu
Ser Leu Gly Trp Gly Tyr Pro Gly Met Gly Asp Phe Ser Tyr Gln Asn
                                        75
Gly Asp Val Glu Lys Glu Ala Asp Val Pro Arg Leu Val Ala Ser Phe
Cys Pro Ser His Pro Pro Thr Lys Asp Met Arg Leu Leu Pro Ser Asn
                                105
            100
Leu Leu Gly Ala Ser Pro Asp Arg Thr Pro Ser Gly Ile
                            120
                                                125
        115
<210> 4869
<211> 418
<212> DNA
<213> Homo sapiens
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tgggaactca atggtgttgc tacctttgga tggactcgga ggcagcccag cttcctggga
120
caggactgca cggactgcct ggggaggggt ctttggcccc ccggttcctg caggggggct
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tqtqagaggc ggggccagag tggccgttgg gaatctgggt gttgcaaggt gaccacaaac
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418
<210> 4870
<211> 125
<212> PRT
<213> Homo sapiens
<400> 4870
Met Ala Met Gly Ile Gly Trp Glu Leu Asn Gly Val Ala Thr Phe Gly
Trp Thr Arg Arg Gln Pro Ser Phe Leu Gly Gln Asp Cys Thr Asp Cys
Leu Gly Arg Gly Leu Trp Pro Pro Gly Ser Cys Arg Gly Ala Arg Gly
Gly Pro Val Ser Ser Trp Ser Gln Val Gly Pro Ile Arg Cys Asp Pro
                        55
Val Pro Pro Gln Gln Pro Trp Arg Arg Gly Thr Leu Pro Ala Val Ala
                                        75
Ala Ala Val Phe Leu Ala Cys Glu Arg Arg Gly Gln Ser Gly Arg Trp
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90
                85
Glu Ser Gly Cys Cys Lys Val Thr Thr Asn Ser Ser Leu Gly Glu Glu
                                105
            100
Glu Glu Asn Ala Ile Asp Phe Gln Glu Pro Ser Glu Val
                            120
                                                125
        115
<210> 4871
<211> 1354
<212> DNA
<213> Homo sapiens
<400> 4871
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120
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gtagetgtgt teattetgga tgtaggetee ggegggtggg ggeaggegag catataeget
420
gagggggaga ctggccgtgg ttcgagaggg gagggctgcc gctctggtga aggctgggcg
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ctaaggccag getecageet eecagetggg gaggeeggea aagtggcagg tgetgaggee
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teettettgg gaagtetgtg gagecacaaa eeegtgagea eeaggetgte cacageeetg
ggetcatget geccaageae eccagagggg aaacgeagae ecaacaegeg eegecaegag
acctecetge gacceegeeg ggtaagcace accgeeeggg cacagaegag geaacggagg
cctcgagaag aaaagcagtt tcctcagcgt catctggcag gtaacagagt ggggcgggtc
1200
caageegget agaetteeeg teeteeeett eeegaetgea tteagteeeg eegggaeegt
1260
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1354
<210> 4872
<211> 90
<212> PRT
<213> Homo sapiens
<400> 4872
Gly Arg Lys Arg Leu Gln Ser Cys Trp Ala Ala Pro Arg Ser Val Gln
Gln Pro Leu Arg Pro Cys Cys Cys Ser Ala Ala Trp Gln Ser Pro Ala
                                25
            20
His Ala Pro Ser Glu Ser Gly Gly His Leu Pro Val Pro Ala Ser Pro
                                                45
        35
Val Pro Ala Pro Ala Ala Ala Trp Ser Val Ser Thr Ala Ala Ala Ala
    50
Pro Ala Ala Cys Arg Pro Ala Ala Gly Ala Gly Pro Cys Gln Gly His
                    70
Gln Gly Leu Pro Gly Ser Pro Leu Pro Glu
                                    90
                85
<210> 4873
<211> 948
<212> DNA
<213> Homo sapiens
<400> 4873
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ccactgtgag ttgaactett tegtgttgae eggecactet cegtgetetg gatgatgteg
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300
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720
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Leu Glu Asn His Val Val Thr Asp Glu Asp Glu Pro Ala Leu Lys Arg
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Gln Arg Leu Glu Ile Asn Cys Gln Asp Pro Ser Ile Lys Ser Phe Leu
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Tyr Ser Ile Asn Gln Thr Ile Cys Leu Arg Leu Asp Ser Ile Glu Ala
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                                        75
Lys Leu Gln Ala Leu Glu Ala Thr Cys Lys Ser Leu Glu Glu Lys Leu
                                     90
Asp Leu Val Thr Asn Lys Gln His Ser Pro Ile Gln Val Pro Met Val
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Ala Gly Ser Pro Leu Arg Thr Thr Gln Met Cys Asn Lys Val Arg Trp
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120
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aaaatacttt gcagctggtg agaaatatca tacctcctct gtcttccaca aagcacaaag
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480
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Val Gly Thr Gly Leu Gly Arg Asp Asp Ala Leu Leu Arg Asn Val Gln
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Gly Ile Leu Glu Val Ser Lys Ala Arg Asp Ile Pro Val Val Ile Asp
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Ala Asp Gly Leu Trp Leu Val Ala Gln Gln Pro Ala Leu Ile His Gly
                    70
                                        75
Tyr Arg Lys Ala Val Leu Thr Pro Asn His Val Glu Phe Ser Arg Leu
                85
                                    90
Tyr Asp Ala Val Leu Arq Gly Pro Met Asp Ser Asp Ser His Gly
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                                105
                                                    110
Ser Val Leu Arg Leu Ser Gln Ala Leu Gly Asn Val Thr Val Val Gln
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                            120
Lys Gly Glu Arg Asp Ile Leu Ser Asn Gly Gln Gln Val Leu Val Cys
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Ser Gln Glu Gly Ser Ser Arg Arg Cys Gly Gly Gln Gly Asp Leu Leu
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Ser Gly Ser Leu Gly Val Leu Val His Trp Ala Leu Leu Ala Gly Pro
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165
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Gln Lys Thr Asn Gly Ser Ser Pro Leu Leu Val Ala Ala Phe Gly Ala
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Cys Ser Leu Thr Arg Gln Cys Asn His Gln Ala Phe Gln Lys His Gly
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Arg Ser Thr Thr Thr Ser Asp Met Ile Ala Glu Val Gly Ala Ala Phe
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Ser Lys Leu Phe Glu Thr
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1140
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                                25
Leu Arg Asp Glu Ser Val Ala His Gly Arg Ile Asp Asn Val Asp Ala
        35
Phe Met Asn Ile Arg Leu Ala Lys Val Thr Tyr Thr Asp Arg Trp Gly
                        55
His Gln Val Lys Leu Asp Asp Leu Phe Val Thr Gly Arg Asn Val Arg
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                    70
Tyr Val His Ile Pro Asp Asp Val Asn Ile Thr Ser Thr Ile Glu Gln
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                                     90
Gln Leu Gln Ile Ile His Arg Val Arg Asn Phe Gly Gly Lys Gly Gln
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Gly Arq Trp Glu Phe Pro Pro Lys Lys Leu
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660
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Ser Leu Ala Ala Ser Ala Gly His Ala Ala Ser Pro Val Leu Pro Ser
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Ala Thr Ala Ser Gly Pro His Val Lys Ser His Leu Thr Arg Val Val
Thr Thr Val Leu Phe Trp Gly Phe Ser Lys Ala Ser Pro Val Val Leu
                                        75
Arg Gly His Ser Glu Gln Ala Asn Thr Ala Arg Val Thr His Tyr Thr
Gln Arg Lys Asp Asn Glu Gln Met Ala Ile Val Glu Asn Ser Val Val
            100
                                105
Cys Phe Ser Asn Ala Thr Tyr Phe Ser Arg Gln Val Ile Leu Pro Met
                            120
Met Thr Ser Ala Thr Lys Leu Arg Ala Arg Gly Leu Pro Met Arg Leu
                        135
                                            140
Val Glu Ser Asn His Val Cys Ser Glu Ala Ser Gly Pro Ser Arg Pro
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145
                    150
Cys His Arg Pro Glu His Arg Thr Val Ile Met Gln Arg Ala Val Thr
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720
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Leu Pro Phe Leu Pro Ser Gln Pro Leu Glv Phe Glv Tvr Met Thr Gln
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Gln Leu Met Asn Leu Ala Gly Gly Ala Val Val Leu Ala Leu Glu Gly
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Gly His Asp Leu Thr Ala Ile Cys Asp Ala Ser Glu Ala Cys Val Ala
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                                         75
Ala Leu Leu Gly Asn Arg Val Ser Arg Leu Pro Pro Pro Ser Met Leu
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Leu Ser Gly Arg
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120
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45

Leu Arg Leu Leu Asn Phe Gln His Asn Phe Ile Thr Arg Ile Gln Asn

40

35

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Leu Leu Gly Lys Asn Arg Ile Lys Lys Ile Ser Asn Leu Glu Asn Leu
Lys Ser Leu Asp Val Leu Asp Leu His Gly Asn Gln Ile Thr Lys Ile
                               105
Glu Asn Ile Asn His Leu Cys Glu Leu Arg Val Leu Asn Leu Ala Arg
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Asn Phe Leu Ser His Val Asp Asn Leu Asn Gly Leu Asp Ser Leu Thr
                       135
Glu Leu Asn Leu Arg His Asn Gln Ile Thr Phe Val Arg Asp Val Asp
                   150
                                       155
Asn Leu Pro Cvs Leu Gln His Leu Phe Leu Ser Phe Asn Asn Ile Ser
               165
                                   170
Ser Phe Asp Ser Val Ser Cys Leu Ala Asp Ser Ser Ser Leu Ser Asp
                               185
Ile Thr Phe Asp Gly Asn Pro Ile Ala Gln Glu Ser Trp Tyr Lys His
                           200
Thr Val Leu Gln Asn Met Met Gln Leu Arg Gln Leu Asp Met Lys Arg
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Ile Thr Glu Glu Glu Arg Arg Met Ala Ser Val Leu Ala Lys Lys Glu
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Glu Glu Lys Lys Arg Glu Ser His Lys Gln Ser Leu Leu Lys Glu Lys
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Lys Arg Leu Thr Ile Asn Asn Val Ala Arg Gln Trp Asp Leu Gln Gln
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Arg Val Ala Asn Ile Ala Thr Asn Glu Asp Arg Lys Asp Ser Asp Ser
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Pro Gln Asp Pro Cys Gln Ile Asp Gly Ser Thr Leu Ser Ala Phe Pro
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Glu Glu Thr Gly Pro Leu Asp Ser Gly Leu Asn Asn Ala Leu Gln Gly
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Leu Ser Val Ile Asp Thr Tyr Leu Val Glu Val Asp Gly Asp Thr Leu
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Ser Leu Tyr Gly Ser Gly Ala Leu Glu Ser Leu Asp Arg Asn Trp Ser
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                               345
Val Gln Thr Ala Gly Met Ile Thr Thr Val Ser Phe Thr Phe Ile Glu
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Phe Asp Glu Ile Val Gln Val Leu Pro Lys Leu Lys Ile Lys Phe Pro
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Asn Ser Leu His Leu Lys Phe Lys Glu Thr Asn Leu Val Met Gln Gln
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<210> 4885
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<sup>&</sup>lt;213> Homo sapiens

<sup>&</sup>lt;400> 4885

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360
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Asn His Pro Asp Ser Ala Ser Glu Lys Asn Pro Val Thr Leu Leu Lys
Glu Leu Ser Val Ile Lys Ser Arg Tyr Gln Thr Leu Tyr Ala Arg Phe
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Lys Pro Val Ala Val Glu Gln Lys Glu Ser Lys Ser Arg Ile Cys Ala
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Ser Ser Trp Asp Tyr Arg Arg Pro Pro Arg Cys Pro Ala Asn Phe Cys
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Ile Phe Ser Lys Asp Arg Val Ser Pro Cys Trp Leu Gly Trp Ser Gln
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Arg Gln Gln Arg Gly Pro Leu Gly Trp Val Gly Val Leu Leu Asp Ser
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Gly Gly Glu His Leu Pro Phe Pro Gln Pro Cys Val His Pro Gln
Met Leu Leu Ala His Arg Ile Ser Gln Cys His Gly Pro Thr Thr Ala
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Ala Lys Ala Ala Ile Val Cys Tyr Asp Leu Thr Asp Ser Ser Ser Phe
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Glu Arq Ala Lys Phe Trp Val Lys Glu Leu Arq Ser Leu Glu Glu Gly
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Cys Gln Ile Tyr Leu Cys Gly Thr Lys Ser Asp Leu Leu Glu Glu Asp
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                                                125
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Ser Ser Cys Ser Ser Ala Leu Trp Gly Val Gly Val Cys Gly Cys Leu
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Asn Lys Gln Thr Ala Val Pro Val Gly Gly Leu Ser Arg Lys Lys Val
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Pro Gln Glu Pro Trp Ala Thr Val Met Glu Lys Arg Leu Gln Glu Ala
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Gln Leu Tyr Lys Glu Glu Gly Asn Gln Arg Tyr Arg Glu Gly Lys Tyr
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                                                 45
Gly Leu Asp His Asn Leu Leu Ala Ser Val Pro Ala Gly Ala Phe Ser
Arg Leu His Lys Leu Ala Arg Leu Asp Met Thr Ser Asn Arg Leu Thr
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Thr Ile Pro Pro Asp Pro Leu Phe Ser Arg Leu Pro Leu Leu Ala Arg
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90
Pro Arg Gly Ser Pro Ala Ser Ala Leu Val Leu Ala Phe Gly Gly Asn
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Pro Leu His Cys Asn Cys Glu Leu Val Trp Leu Arg Arg Leu Ala Arg
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Phe Thr Val Ser Glv Ile Gln Val Arg Tyr Met Lys Ile Ile Glu Lys
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-		275				-	280		_			285			
Ser	Dro		Δla	Len	Ser	Ser		Pro	Asn	Asn	Len		Pro	Thr	Glv
DCI	290	OCI		204	001	295	001				300				
TT week		C1 n	Dwo	Tara	Thr		17-1	Dro	71-	Gln.		C1.,	7 200	719	Dro
	Ser	GIII	PIO	Буб	310	PIO	vai	FIO	мта	315	Arg	GIU	Arg	AIA	320
305						_	_	_			_	_			
Val	Ser	GIY	Thr		Glu	ьys	Asn	Lys		Arg	Pro	Arg	GIY		Arg
				325					330					335	
Asp	Ser	Ser		Tyr	Trp	Glu	Ile		Ala	Ser	Glu	Val		Leu	Ser
			340					345					350		
Thr	Arg	Ile	Gly	Ser	Gly	Ser	Phe	Gly	Thr	Val	Tyr	Lys	Gly	Lys	Trp
		355					360					365			
His	Gly	Asp	Val	Ala	Val	Lys	Ile	Leu	Lys	Val	Val	Asp	Pro	Thr	Pro
	370	-				375			-		380	_			
Glu	Gln	Phe	Gln	Ala	Phe	Ara	Asn	Glu	Val	Ala	Val	Leu	Arq	Lvs	Thr
385					390	3				395					400
	Hic	Va 1	a en	Tle	Leu	T.011	Phe	Met	Glv		Met	Thr	Lvs	Asn	
ALG	1113	vai	Abii	405	Бец	Бец	FILE	Hec	410	TYL	PIC C	1111	Буз	415	no
		-1-	**- 1		a1-	m		G1				T	m		77.6
Leu	AIA	TIE		Int	Gln	пр	cys		GIY	ser	ser	Leu		Lys	nis
		_	420			_		425				_	430	_	
Leu	His		Gln	Glu	Thr	Lys		Gln	Met	Phe	Gln		Ile	Asp	IIe
		435					440					445			
Ala		Gln	Thr	Ala	Gln		Met	Asp	Tyr	Leu		Ala	Lys	Asn	Ile
	450					455					460				
Ile	His	Arg	Asp	Met	Lys	Ser	Asn	Asn	Ile	Phe	Leu	His	Glu	Gly	Leu
465					470					475					480
Thr	Val	Lys	Ile	Gly	Asp	Phe	Gly	Leu	Ala	Thr	Val	Lys	Ser	Arg	Trp
		-		485			-		490					495	
Ser	Glv	Ser	Gln	Gln	Val	Glu	Gln	Pro	Thr	Gly	Ser	Val	Leu	Trp	Met
	•		500					505		•			510	-	
Ala	Pro	Glu		He	Arg	Met	Gln		Asn	Asn	Pro	Phe	Ser	Phe	Gln
		515			5		520					525			
car	Acn		Tur	Car	Tyr	Glv		Wal.	T.011	Tur	Glu		Mot	Thr	Glv
Ser	530	VAL	- Y -	Ser	- 7 -	535	110	vai	Deu	- 7 -	540	Deu	rice	1111	017
a1		D	m		His			7	3	7		T1 .	T1 .	Dh a	Mak
	Leu	PIO	ıyr	ser		TIE	ASII	ASII	Arg		GIII	116	ire	Pne	
545		_		_	550	_	_	_		555	_	_	_		560
Val	Gly	Arg	Gly		Ala	Ser	Pro	Asp		Ser	Lys	Leu	Tyr		Asn
				565					570					575	
Cys	Pro	Lys		Met	Lys	Arg	Leu		Ala	Asp	Cys	Val		Lys	Val
			580					585					590		
Lys	Glu	Glu	Arg	Pro	Leu	Phe	Pro	Gln	Ile	Leu	Ser	Ser	Ile	Glu	Leu
		595					600					605			
Leu	Gln	His	Ser	Leu	Pro	Lys	Ile	Asn	Arg	Ser	Ala	Ser	Glu	Pro	Ser
	610					615			_		620				
Lev		Arc	Ala	Ala	His		Glii	Asp	Ile	Asn		Cvs	Thr	Lev	Thr
625					630					635		-1-			640
	80~	Drc	Ar~	T.011	Pro	1721	Dhe			033					
TILL	JCI	0	9	645	-10	vul	- 110								
				043											

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tacccqtatq cccatctctc aqctqaqqac tttaatatct atqqccatqq qqqccqccaq
ttetggetgg teageteetg ettettette etgeteggag gagettetae gtgtatgegg
gcatcctggc accgctcaac n
261
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<212> PRT
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Ile Ile Leu Gly Leu Ala Phe Gly Xaa Leu Glu Ser Lys Ser Ser Ile
Lvs Arg Val Leu Ala Ile Thr Thr Val Leu Ser Pro Ala Leu Ser Val
                                25
Thr Gln Gly Thr Arg Lys Ile Leu Tyr Pro Tyr Ala His Leu Ser Ala
Glu Asp Phe Asn Ile Tyr Gly His Gly Gly Arg Gln Phe Trp Leu Val
                        55
Ser Ser Cys Phe Phe Phe Leu Leu Gly Gly Ala Ser Thr Cys Met Arg
                                        75
                                                             80
Ala Ser Trp His Arg Ser Thr
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ccaagggctg ggcatggcgg caccgctggt tcaccctctc tcgtcttcct ccacaggtgt
getteeegea cagetgeage catggggtet gaggaceagg gegeecagaa ceccagetgt
aaaatcatqa cqtttcqccc aaccatqqaa qaatttaaaq acttcaacaa atacqtqqcc
tacatagagt cgcagggagc ccaccgggcg ggcctggcca agatcatccc cccgaaggag
360
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tggaagccgc ggcagacgta tgatgacatc gacgacgtgg tgatcccggc gcccatccag
caggtggtga cgggccagtc gggcctcttc acgcagtaca atatccagaa gaaggccatg
480
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gactttgacg accttgaacg caaatactgg aagaacctca cctttgtctc cccgatctac
qqqqctqaca tcaqcqqctc tttqtatqat qacqtaaqta tqaqqctccq qqqaaqaaca
gggaccaget teetggtggg tggtggtggg agggeeetga aegggaetet geettggeag
720
atquagette caqqeaqqea aqqttaacce ceteqeecaq qetetqqatq eqqqeeteqe
cctqtqqtqa cqaaaqaqqa aqccaqqctt tctctqattt ttqcaqqqcc cctcctqcct
840
caccetgeaq ecceaceet gaqeteacee tqqeeccace tetqqeetea geageeggee
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cacagogtgt tacaaacacg tqtactttcc cagtccctgc cgctcgtctt cctggcactg
tggageeteg agtee
975
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Thr Phe Arg Pro Thr Met Glu Glu Phe Lys Asp Phe Asn Lys Tyr Val
            20
                                25
                                                    30
Ala Tyr Ile Glu Ser Gln Gly Ala His Arg Ala Gly Leu Ala Lys Ile
Ile Pro Pro Lys Glu Trp Lys Pro Arg Gln Thr Tyr Asp Asp Ile Asp
Asp Val Val Ile Pro Ala Pro Ile Gln Gln Val Val Thr Gly Gln Ser
Gly Leu Phe Thr Gln Tyr Asn Ile Gln Lys Lys Ala Met Thr Val Gly
Glu Tyr Arg Arg Leu Ala Asn Ser Glu Lys Tyr Cys Thr Pro Arg His
Gln Asp Phe Asp Asp Leu Glu Arg Lys Tyr Trp Lys Asn Leu Thr Phe
                            120
Val Ser Pro Ile Tyr Gly Ala Asp Ile Ser Gly Ser Leu Tyr Asp Asp
    130
Val Ser Met Arg Leu Arg Gly Arg Thr Gly Thr Ser Phe Leu Val Gly
145
                    150
                                        155
                                                             160
Gly Gly Gly Arg Ala Leu Asn Gly Thr Leu Pro Trp Gln Met Lys Leu
                                                         175
                165
                                    170
Pro Gly Arg Gln Gly
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ccatqaattq tcatttatag tccaattttt tatcttaatc ataaaatgtt taggaatcta
tqaaatttaa ctttaggaac aaaacgttta gcagggttga ttgatattat ttttacattg
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Asn Phe Asp Ser Val Glu Leu His Gly Thr Met Lys Ser Tyr Phe Gly
Gly Leu Leu Cys Val Cys Trp Ser Pro Asp Gly Lys Tyr Ile Val Thr
                            40
Gly Gly Glu Asp Asp Leu Val Thr Val Trp Ser Phe Val Asp Cys Arg
                        55
Val Ile Ala Arq Gly His Gly His Lys Ser Trp Val Ser Val Val Ala
                                        75
Phe Asp Pro Tyr Thr Thr Ser Val Glu Glu Gly Asp Pro Met Glu Phe
Ser Gly Ser Asp Glu Asp Phe Gln Asp Leu Leu His Phe Gly Glu Ile
                                105
Glu Gln Ile Val His Ser Pro Gly Ser Pro Asn Gly Thr Leu Gln Thr
                            120
                                                125
Ala Ala Pro Ser Val Thr Tyr Arg Phe Gly Ser Val Gly Gln Asp Thr
                        135
                                            140
Gln Leu Cys Leu Trp Asp Leu Thr Glu Asp Ile Leu Phe Pro His Gln
                    150
                                        155
Pro Leu Ser Arg Ala Arg Thr His Thr Asn Val Met Asn Ala Thr Ser
                165
                                    170
Pro Pro Ala Gly Ser Asn Gly Asn Ser Val Thr Thr Pro Gly Asn Ser
                                                    190
            180
                                185
Val Pro Pro Pro Leu Pro Arg Ser Asn Ser Leu Pro His Ser Ala Val
                            200
                                                205
Ser Asn Ala Gly Ser Lys Ser Ser Val Met Asp Gly Ala Ile Ala Ser
                        215
                                            220
Gly Val Ser Lys Phe Ala Thr Leu Ser Leu His Asp Arg Lys Glu Arg
                    230
                                        235
His His Glu Lys Asp His Lys Arg Asn His Ser Met Gly His Ile Ser
                245
                                    250
Ser Lys Ser Ser Asp Lys Leu Asn Leu Val Thr Lys Thr Lys Thr Asp
            260
                                265
Pro Ala Lys Thr Leu Gly Thr Pro Leu Cys Pro Arg Met Glu Asp Val
                            280
                                                285
Pro Leu Leu Glu Pro Leu Ile Cys Lys Lys Ile Ala His Glu Arg Leu
                        295
                                            300
Thr Val Leu Ile Phe Leu Glu Asp Cys Ile Val Thr Ala Cys Gln Glu
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305
                    310
                                        315
Gly Phe Ile Cys Thr Trp Gly Arg Pro Gly Lys Val Val Ser Phe Asn
                325
                                    330
                                                        335
Pro
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aaqqqqqqq teccacacte tetqqqteca qqcacaaaqc tatcetecgt tgttetgate
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caaqqaqqcc qccaqccqqq ctqtaccctt accttggggg tgtgtgcaga tggaaggtgg
qaaqaqacaq accaacaqqa agtgttctct tcaggggttg ccagccccac cctgaatctc
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420
cagagggagg cacccgagct ggtgtgagca gctacgtggg gtggtggtcc agggaacaga
gggagggcac tggagccatt gcctgcctag ttcagtcctc aaatgggtcc aagccagctc
540
aggtctgcag cgccaggccc agggtacctg gcagcccagc cgatggtacc attggctggt
geteceactq aggtettqaq aaggtaatqq qqaqaqccac ttgcccctgc ctctgtcccc
aqtqqacttc tttttgttca aggccaaatg ccaccccgtc agagagagga ccggt
715
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Met Lvs Arg Gly Val Pro His Ser Leu Gly Pro Gly Thr Lys Leu Ser
                                    10
Ser Val Val Leu Ile Cys Arg Ala Ser Ala Leu Ser Arg Tyr Leu Val
Val Ala Glu Pro Trp Pro Thr Arg Ser Gln Gly Gly Arg Gln Pro Gly
        35
Cys Thr Leu Thr Leu Gly Val Cys Ala Asp Gly Arg Trp Glu Glu Thr
Asp Gln Glu Val Phe Ser Ser Gly Val Ala Ser Pro Thr Leu Asn
                    70
Leu Arg Ala Ser Ser Pro Ala Lys Ala Arg Ala Leu Ser Arg Pro
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95
                85
                                     90
Trp Ala Leu Tyr Lys Gln Arg Glu Ala Pro Glu Leu Val
            100
                                105
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teggeeteta eeegeeetee eeaaggteet eeeteeetgg acteaaaage etetaettgg
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quatquete ctqctqctqc ctcttccttt tctttcgagt cccagccttg cccaagcgcc
300
cettecaaaq etteaceaqe gecageageg etgatgtgtg ggaccacate acceeccata
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gettgeagee tecageetet tetggatgtt etgteageet eegeeteete ateeteagtt
480
tetetggeat aggeetetee cagtgaeggg caaggeeetg egtetgeece tgtgetteeg
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730
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Ser Arg Ser Pro Pro Ala Ser Thr Met Ala Pro Ile Pro Ser Ala Leu
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Ala Val Trp Glu Pro Ala Gly Ser Ser Pro Gln Leu Ser Ser Ala Pro
            20
                                25
Ala Asp Ser Ser Ala Ser Thr Arg Pro Pro Gln Gly Pro Pro Ser Leu
        35
                                                 45
Asp Ser Lys Ala Ser Thr Trp Leu Pro Leu Pro Val Thr Ser Ser Ser
                                             60
                        55
Ala Glu Pro Ser Arg Pro Asn Ser Cys Pro Pro Ala Cys Ser Pro Ala
                                        75
                    70
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Ala Ala Ser Ser Phe Ser Phe Glu Ser Gln Pro Cys Pro Ser Ala Pro

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90
Ser Lys Ala Ser Pro Ala Pro Ala Ala Leu Met Cys Gly Thr Thr Ser
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                                105
                                                     110
Pro Pro Ile Ile Pro Ala Ala Thr Glu Pro Val Cys Ala Ser Ser Arg
                            120
                                                 125
        115
Ser Gly Arg Pro Thr Ala Thr Ala Cys Ser Leu Gln Pro Leu Leu Asp
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Val Leu Ser Ala Ser Ala Ser Ser Ser Ser Val Ser Leu Ala
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                                        155
145
<210> 4941
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gatetgetee tgggcageet gaaggagaag eeegteacca aggagggeeg ggettecate
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1020
gcaccacgca ctcaggccat ggaggagcag ctggtcagca ccttggtgcc cctactgctg
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1140
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cgtctgaatg aagtgaaagc tgctctggat aacttgagac atgacccaga agcatcagtg
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Pro Pro Lys Asp Thr Lys Lys Gly Ala Gln Pro Ser Pro Phe Val Pro
        35
                            40
Val Arg Trp Val Val Lys Val Lys Thr Leu Leu Arg Met Gly
                        55
                                            60
Cys Ser Tyr Glu Thr Thr Phe Leu Glu Asp Gln Gly Gly Trp Glu Leu
65
                    70
                                        75
Met Glu Gln Val Glu Ser His His Arg Gly Val Ala Leu Leu Ala Arg
                85
                                    90
Ala Met Val Gln Tyr Ser Cys Gln Glu Leu Cys Arg Ile Leu Tyr Leu
            100
                                105
Leu Ile Pro Leu Leu Glu Arg Gly Asp Glu Lys His Arg Ile Thr Ala
                            120
                                                125
Thr Ala Phe Phe Val Glu Leu Leu Gln Met Glu Gln Val Arg Arg Ile
                        135
Pro Glu Glu Tyr Ser Leu Gly Arg Met Ala Glu Gly Leu Ser His His
                    150
                                        155
Asp Pro Ile Met Lys Val Leu Ser Ile Arg Gly Leu Val Ile Leu Ala
                                    170
Arg Arg Ser Glu Lys Thr Ala Lys Val Lys Ala Leu Leu Pro Ser Met
                                185
                                                    190
Val Lys Gly Leu Lys Asn Met Asp Gly Met Leu Val Val Glu Ala Val
                            200
His Asn Leu Lys Ala Val Phe Lys Gly Arg Asp Gln Lys Leu Met Asp
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Ser Ala Val Tyr Val Glu Met Leu Gln Ile Leu Leu Pro His Phe Ser
225
                    230
                                        235
Asp Ala Arg Glu Val Val Arg Ser Ser Cys Ile Asn Leu Tyr Gly Lys
                                    250
                245
Val Val Gln Lys Leu Arg Ala Pro Arg Thr Gln Ala Met Glu Gln Gln
            260
                                265
Leu Val Ser Thr Leu Val Pro Leu Leu Leu Thr Met Gln Glu Gly Asn
                            280
        275
Ser Lys Val Ser Gln Lys Cys Val Lys Thr Leu Leu Arg Cys Ser Tyr
                        295
                                             300
Phe Met Ala Trp Glu Leu Pro Lys Arg Ala Tyr Ser Arg Lys Pro Trp
                    310
                                        315
Asp Asn Gln Gln Gln Thr Val Ala Lys Ile Cys Lys Cys Leu Val Asn
                325
                                    330
Thr His Arg Asp Ser Ala Phe Ile Phe Leu Ser Gln Ser Leu Glu Tyr
                                345
            340
Ala Lys Asn Ser Arg Ala Ser Leu Arg Lys Cys Ser Val Met Phe Ile
                            360
                                                 365
        355
Gly Ser Leu Val Pro Cys Met Glu Ser Ile Met Thr Glu Asp Arg Leu
                        375
                                             380
Asn Glu Val Lys Ala Ala Leu Asp Asn Leu Arg His Asp Pro Glu Ala
                    390
                                        395
Ser Val Cys Ile Tyr Ala Ala Gln Val Gln Asp His Ile Leu Ala Ser
                405
                                     410
                                                         415
Cys Trp Gln Asn Ser Trp Leu Pro His Gly Asn Ser Trp Val Cys Tyr
                                                     430
                                 425
Ser Ala Thr Thr His Arg Trp Ser Pro Ser Cys Glu Asn Leu Pro Thr
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Ser His Gln Arg Arg Ser Trp Ile Met Gln Ala Leu Gly Ser Trp Lys
                                             460
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Met Ser Leu Lys Lys
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tgccqtaatt tcctctcagg cgcaattact ctcttccata ttggttaaca gtagaaggct
cagtttetet geteateaca eggeettegg caetgtaget ttgggtggtg ggetgeagat
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taattttqta accaccttaa gaaaaatacg gaactctaac teettgecac tcaagaaatg

420

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tectecettt cagaatatge etteegeatg tetegtetea gtgeeegget atttggtgaa
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qaqaaqaaq tqcatttatt qtctttccac atattqqaqq aatgtcatct tcctaaatga
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aqtttatttq qaqqaacaca qtcatctcct tqqtqaaatc taatccgqtt acattqtggc
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<212> PRT
<213> Homo sapiens
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Arg Leu Phe Gly Glu Val Thr Arg Pro Thr Asn Ser Lys Ser Met Lys
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Val Val Lys Leu Phe Ser Glu Leu Pro Leu Ala Lys Lys Glu Thr
Tyr Asp Trp Tyr Pro Asn His His Thr Tyr Ala Glu Leu Met Gln Thr
Leu Arg Phe Leu Gly Leu Tyr Arg Asp Glu His Gln Asp Phe Met Asp
65
Glu Gln Lvs Arg Leu Lvs Lvs Leu Arg Glv Lvs Glu Lvs Pro Lvs Lvs
Glv Glu Glv Lvs Arg Ala Ala Lvs Arg Lvs
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<210> 4945
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<212> DNA
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tacaacacat atgatgtcca cttttatgct teetttgccc teatcatgct ctggcccaaa
180
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cttgagetea geetacagta tgacatgget etggeeaete teagggagga eetgacaegg cgacggtacc tgatgagtgg ggtgatggca cctgtgaaaa ggaggaacgt catcccccat gatattgggg acccagatga tgaaccatgg ctccgcgtca atgcatattt aatccatgat actgctgatt ggaaggacct gaacctgaag tttgtgctgc aggtttatcg ggactattac ctcacgggtg atcaaaactt cctgaaggac atgtggcctg tgtgtctagt aagggatgca catgcagtgg ccagtgtgcc aggggtatgg ttggtgtctg ggaagagcct agctggttgt tgeettteet eggtaeetag gtetteaaca tettggteee tetetagget gtgatggaat ctgaaatgaa gtttgacaag gaccatgatg gactcattga aaatggaggc tatgcagacc agacctatga tggatgggtg accacaggcc ccaggttagc gggtaggggt ttccaggagg cctgaggtga gaaactgggc aacaagggat tgtagggctc aagaaagaat gactcattgt ctattacacg gcatgggage agetggaget gccagtetga cccccaaacc catgtccctg atcagtgctt actgtggagg gctgtggctg gcagctgtgg ctgtgatggt ccagatggct getetgtgtg gggeacagga catecaggat aagttttett etateeteag eeggggeeaa gaageetatg agagaetget gtggaatggt gagttegggg ageetaagta gtettaagge 1020 agetgagagg acaccaggag cettattttt etetteeteg actecaggee getattacaa 1080 ctatgacage agetetegge etcagteteg tagtgttatg tetgaccagt gtgctggaca gtggttcctg aaggcctgtg gcctaggaga aggagacact gaggtgtttc ctacccaaca tgtggtccgt gctctccaaa ctatctttga gctgaacgtc caggcctttg caggaggggc 1260 catgggggct gtgaatggga tgcagccca tggtgtccct gataaatcca gtgtgcagtc tgatgaagtc tgggtgggtg tggtctacgg gctggcagct accatgatcc aagaggtaat geacteettt teecatetet ceaceatetg tateetggee cagaaaactt ceteaaceae caaatttctt caaggcataa cccaatgcca tcttgtccgt ctataaagcc tcccattttt ccctggtatg cattccaget cctgccttca ggcttctgtc tgtgggtcat agttatctcc tccacttgct gggagetect tgaaggcaaa gactctactg cetecateta tecagtggaa 1620 gtggetette agagggtgee aagttagtat gtatgactgt cateteteec aacagggeet 1680 gacttgggag ggcttccaga cagctgaagg ctgctaccgt accgtgtggg agcgcctggg 1740 tetggeette cagaccccag aggeatactg ccagcagega gtgtteegeg eg 1792

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<212> PRT
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Pro Pro Gly Gln Glu Tyr Arg Met Tyr Asn Thr Tyr Asp Val His Phe
Tyr Ala Ser Phe Ala Leu Ile Met Leu Trp Pro Lys Leu Glu Leu Ser
                        55
Leu Gln Tyr Asp Met Ala Leu Ala Thr Leu Arg Glu Asp Leu Thr Arg
                    70
                                        75
Arg Arg Tyr Leu Met Ser Gly Val Met Ala Pro Val Lys Arg Arg Asn
                                    90
                                                         95
Val Ile Pro His Asp Ile Gly Asp Pro Asp Asp Glu Pro Trp Leu Arg
            100
                                105
Val Asn Ala Tyr Leu Ile His Asp Thr Ala Asp Trp Lys Asp Leu Asn
                            120
                                                 125
Leu Lys Phe Val Leu Gln Val Tyr Arg Asp Tyr Tyr Leu Thr Gly Asp
                        135
                                            140
Gln Asn Phe Leu Lys Asp Met Trp Pro Val Cys Leu Val Arg Asp Ala
                    150
His Ala Val Ala Ser Val Pro Gly Val Trp Leu Val Ser Gly Lys Ser
                                    170
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Leu Ala Gly Cys Cys Leu Ser Ser Val Pro Arg Ser Ser Thr Ser Trp
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                                                     190
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Ser Leu Ser Arg Leu
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cacttqaata acaqccctqt ggcattttag atctcqagca ctgggatttg tcaattgtca
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tqqaaqacta ctaaqcacqt agtttcaqtc attcagttqa tagacatttg aacacttatg
420
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Met Leu Pro His Asn Ile Pro Ser Ser Leu Ser Leu Leu Thr Ser Met
            20
Val Asp Asp Met Trp His Tyr Ala Gly Asp Gln Ser Thr Asp Phe Asn
                                                 45
Trp Tyr Thr Arg Arg Ala Met Leu Ala Ala Ile Tyr Asn Thr Thr Glu
                        55
                                             60
    50
Leu Val Met Met Gln Asp Ser Ser Pro Asp Phe Glu Asp Thr Trp Arg
                                        75
65
Phe Leu Glu Asn Arg Val Asn Asp Ala Met Asn Met Gly His Thr Ala
                                     90
Lys Gln Val Lys Ser Thr Gly Glu Ala Leu Val Gln Gly Leu Met Gly
            100
                                105
Ala Ala Val Thr Leu Lys Asn Leu Thr Xaa Leu Asn Gln Arg Arg
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                            120
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gettgggagg aaaagacget gtecaagtac gagtecageg agattegeet getggagate
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300
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360
gcatgccagg gcggatccca gaggccctgc agcgggaatg gccactgcag cggagatggg
420
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qactqcatqq acqgctactt cagctcqctc cqgaacgaqa cccacagcat ctgcacagcc
tgtgacgagt cctgcaagac gtgctcgggc ctgaccaaca gagactgcgg cgagtgtgaa
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660
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cetecetgea gegetgegea gttetgtaag aacgecaacg geteetacae gtgegaagag
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tctqqctacq cqagqgagca cggacagtgt gcagatgtgg acgagtgctc actagcagaa
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taccetttaa attatteaga aggatgteee gtggaaaatg tggeeetgag gatgeegtet
cctgcagtgg acagcggcgg ggagaggctg cctgctctct aacggttgat tctcatttgt
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Arg Gly Leu Val Asp Lys Phe Asn Gln Gly Met Val Asp Thr Ala Lys
Lys Asn Phe Gly Gly Gly Asn Thr Ala Trp Glu Glu Lys Thr Leu Ser
                           40
                                               45
Lys Tyr Glu Ser Ser Glu Ile Arg Leu Leu Glu Ile Leu Glu Gly Leu
Cys Glu Ser Ser Asp Phe Glu Cys Asn Gln Met Leu Glu Ala Gln Glu
                   70
Glu His Leu Glu Ala Trp Trp Leu Gln Leu Lys Ser Glu Tyr Pro Asp
Leu Phe Glu Trp Phe Cys Val Lys Thr Leu Lys Val Cys Cys Ser Pro
           100
Gly Thr Tyr Gly Pro Asp Cys Leu Ala Cys Gln Gly Gly Ser Gln Arg
Pro Cys Ser Gly Asn Gly His Cys Ser Gly Asp Gly Ser Arg Gln Gly
                       135
                                           140
Asp Gly Ser Cys Arg Cys His Met Gly Tyr Gln Gly Pro Leu Cys Thr
                   150
                                       155
145
Asp Cys Met Asp Gly Tyr Phe Ser Ser Leu Arg Asn Glu Thr His Ser
               165
                                   170
Ile Cys Thr Ala Cys Asp Glu Ser Cys Lys Thr Cys Ser Gly Leu Thr
                                                   190
                               185
Asn Arg Asp Cys Gly Glu Cys Glu Val Gly Trp Val Leu Asp Glu Gly
                           200
                                               205
Ala Cys Val Asp Val Asp Glu Cys Ala Ala Glu Pro Pro Pro Cys Ser
```

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210
                        215
                                            220
Ala Ala Gln Phe Cys Lys Asn Ala Asn Gly Ser Tyr Thr Cys Glu Glu
                                        235
225
Cys Asp Ser Ser Cys Val Gly Cys Thr Gly Glu Gly Pro Gly Asn Cys
                245
                                    250
Lys Glu Cys Ile Ser Gly Tyr Ala Arg Glu His Gly Gln Cys Ala Asp
                                265
Val Asp Glu Cys Ser Leu Ala Glu Lys Thr Cys Val Arg Lys Asn Glu
        275
                            280
Asn Cys Tyr Asn Thr Pro Gly Ser Tyr Val Cys Val Cys Pro Asp Gly
Phe Glu Glu Xaa Gly Arg Cys Leu Cys Ala Ala Gly Arg Gly
                    310
305
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<211> 1835
<212> DNA
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900
cagtecactg attitaactg gtacaccege egagecatge tggetgecat etacaacaca
acaqagetgg tgatgatgca ggactectet ecagactttg aggacacttg gegetteetg
1020
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gaaaaccggg ttaatgatgc aatgaacatg ggccacactg ccaagcaggt aaagtccaca
ggagaggcac tggtgcaagg actcatgggt gcagcagtga cgctcaagaa cttgacaggt
1140
ctaaaccagc gtcggtgaga ggaaggggta taagctacaa tgcctagaag agaatgagcg
gacagattga aagagetttg aaaagtataa ggtgccatcc acataacctg gtgttcacga
qaacacacta aaqqactcct gagtcactac cacagccacc tggaaaccac aaggcatttg
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1440
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tgagacaagt gcctgctgga cagaggtgtg attccaggcc tggtgtcaca tgacaccagc
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                                25
            20
Val Pro Arg Ala Phe His Ala Ser Ala Val Gly Leu Arg Ser Ser Asp
                            40
Glu Gln Lys Gln Gln Pro Pro Asn Ser Phe Ser Gln Gln His Ser Glu
Thr Gln Gly Ala Glu Lys Pro Asp Pro Glu Ser Ser His Ser Pro Pro
                    70
                                        75
Arg Tyr Thr Asp Gln Gly Gly Glu Glu Glu Asp Tyr Glu Ser Glu
                                    90
Glu Gln Leu Gln His Arg Ile Leu Thr Ala Ala Leu Glu Phe Val Pro
                                105
                                                     110
            100
Ala His Gly Trp Thr Ala Glu Ala Ile Ala Glu Gly Ala Gln Ser Leu
                                                 125
                            120
        115
Gly Leu Ser Ser Ala Ala Ala Ser Met Phe Gly Arg Met Gly Ser Glu
                                            140
                        135
Leu Ile Leu His Phe Val Thr Gln Cys Asn Thr Arg Leu Thr Arg Val
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145
                    150
                                        155
Leu Glu Glu Glu Gln Lys Leu Val Gln Leu Gly Gln Ala Glu Lys Arg
                165
                                    170
Lys Thr Asp Gln Phe Leu Arg Asp Ala Val Glu Thr Arg Leu Arg Met
                                185
            180
Leu Ile Pro Tyr Ile Glu His Trp Pro Arg Ala Leu Ser Ile Leu Met
                            200
Leu Pro His Asn Ile Pro Ser Ser Leu Ser Leu Leu Thr Ser Met Val
    210
                        215
Asp Asp Met Trp His Tyr Ala Gly Asp Gln Ser Thr Asp Phe Asn Trp
                    230
                                        235
Tyr Thr Arg Arg Ala Met Leu Ala Ala Ile Tyr Asn Thr Thr Glu Leu
                245
                                    250
Val Met Met Gln Asp Ser Ser Pro Asp Phe Glu Asp Thr Trp Arg Phe
            260
                                265
Leu Glu Asn Arg Val Asn Asp Ala Met Asn Met Gly His Thr Ala Lys
        275
                            280
                                                285
Gln Val Lys Ser Thr Gly Glu Ala Leu Val Gln Gly Leu Met Gly Ala
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                        295
                                            300
Ala Val Thr Leu Lys Asn Leu Thr Gly Leu Asn Gln Arg Arg
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<212> DNA
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240
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<213> Homo sapiens
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Ser Ala Trp Gly Cys Leu Ala Ala Ser Pro Val Leu Gly Ala Gly Ile
            20
                                25
Thr Trp Pro Arg Val Pro Pro Gly Gly Ser Leu Lys Glu Gly Arg Ala
Val Gly Arg Ser Gln Arg Gly Pro Thr Pro Gln Asn Ala His Lys Ser
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50
                        55
                                             60
Trp Asn Gln Leu Val Thr Ala Ala Gly Pro Ser Arg Pro Ile Trp Ile
                                                             80
Asp Pro Leu Gly Thr His Cys Thr Arg Glu Pro Gln Met Gln Leu Ser
Ser Met Gly Gly Ala Leu Ser Ala Gly Gly Val Trp Asp Arg Arg Arg
                                105
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Glu Ala
<210> 4955
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<212> DNA
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ageteageet geeeaggaac aactetggge aagagatgtg gaaagaaaga geteangggg
gggcacgcat ggcatcctgg ggggacatct gagggcaccc ccacccacta ttcctccctc
caaggtggcc totgagtgtg aaggcagggg gaagcagaca cotgcccctc actotccctc
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364
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<213> Homo sapiens
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Met Gly Thr Glu His Leu Gly Leu Arg Pro Glu Glu Gln Thr Ala Arg
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Gln Gly Gly Arg Gly His Gln Pro Pro Pro Phe Cys Asp Ile Arg Thr
Arg Ala Gln Pro Ala Gln Glu Gln Leu Trp Ala Arg Asp Val Glu Arg
                            40
Lys Ser Ser Xaa Gly Gly Thr His Gly Ile Leu Gly Gly His Leu Arg
                        55
Ala Pro Pro Pro Thr Ile Pro Pro Ser Lys Val Ala Ser Glu Cys Glu
                    70
                                        75
                                                             80
Gly Arg Gly Lys Gln Thr Pro Ala Pro His Ser Pro Ser Leu Pro His
Ser Tyr Arg Val Gly Gly Val Pro Gly Met Ile Pro Glu Gly Arg Ile
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                                                     110
Gln Gly
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<211> 872
<212> DNA
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aacccacagc acctcctgca gtcctggagg gaaaagggac agtaacatga agtgtctgaa
gatecattte acctettte catqtqaate atqacqettt caatqcattt ettqacaqqa
300
ttctattttq aaaqaatqat qctcaatctq taccttttat qcttcttqtt tcttctccat
360
caataatatg tcagtcaact gcttgtcaga gacacttagc tgctgacagg tcctcataac
420
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tgagaatgca aatgcaggac atqaacagta atgacaagaa qccaaacatg tgtatgtttt
540
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qqactqtcqa tcaactaqqt ccaaqqcctq qqtqqctqat qaqccaaaqa qaaacttcaq
cgataacaga tattcatcag gaattcggtc ccgtacttcg cgcgctctcc tgcaccgccg
cegecatete geteaggage tectecacaa cegeeggeaa etaeggeeat egegeegeag
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gacacgccct ccacgacgcg gaccgcgcga cgctccagct gactgcgcct acctgtggag
840
gatectgace eccegeegge etegtteega at
872
<210> 4958
<211> 51
<212> PRT
<213> Homo sapiens
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Gln Ile Phe Ile Arg Asn Ser Val Pro Tyr Phe Ala Arg Ser Pro Ala
Pro Pro Pro Pro Ser Arg Ser Gly Ala Pro Pro Gln Pro Pro Ala Thr
            20
                                25
Thr Ala Ile Ala Pro Gln Asp Thr Pro Ser Thr Thr Arg Thr Ala Arg
        35
                            40
                                                 45
Arg Ser Ser
    50
<210> 4959
<211> 449
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<212> DNA
<213> Homo sapiens
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	_	Leu 275					280			_	-	285			
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Glu Gly Ala Arg Pro Gly Gly Ile Ile His Val Tyr Gly Asp Asp Ser
Ser Asp Arg Ala Ala Ser Ser Phe Ile Pro Tyr Cys Ser Met Ala Gln
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Ser Lys Leu Gln Val Asn Leu Leu Ser Lys Phe Xaa Leu Ile Ala Lys
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		D	34U	Tyr	) cn	Tla	Glv		Tle	Va 1	Glu	Phe			Asp
ser	Asp			TYL	Mon	116	360					365		•	-
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Pro			ser	Leu	GIU	375	GIY	FIO	7110		380		-2		
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			_	405	**- 1	a1	<b>~1</b>	C1.,			Tla	Trn	Lvs		His
Trp	Pro	GIU			vai	GIU	GIY	425	Map	Cys	110		430		
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545

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Val Gly Pro Pro Phe Leu Met Asp Glu Asn Ser Trp Phe Asn Lys Cys
Lys Arg Val Lys Gln Lys Tyr Gln Leu Thr Leu Glu Gln Lys Gly Tyr
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Ser Gln Asn Lys Ile Leu Leu Gln Arg Ile Glu Asp Ser Asp Leu Ala
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His Lys Leu Glu Lys Glu Gln Leu Glu Tyr Ile Ile Val Glu Leu Gln
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Asp Gln Leu Thr Val Leu Lys Asn Asn Asp Leu Arg Ser Arg Gln Glu
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Ala Val Pro Ala Ser Leu Val Gln Pro Gly Val Leu Arg Cys Tyr Cys
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Pro Ala His Glu Val Gly Leu Val Ser Leu Gln Val Ala Gly Arg Glu
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Gly Pro Leu Ser Ala Ser Val Leu Phe Glu Tyr Arg Ala Arg Arg Phe
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                                    650
Leu Ser Leu Pro Ser Thr Gln Leu Asp Trp Leu Ser Leu Asp Asp Asn
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Gln Phe Arg Met Ser Ile Leu Glu Arg Leu Glu Gln Met Glu Lys Arg
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Met Ala Glu Ile Ala Ala Ala Gly Gln Val Pro Cys Gln Gly Pro Asp
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Ala Pro Pro Val Gln Asp Glu Gly Gln Gly Pro Gly Phe Glu Ala Arg
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                                        715
Val Val Val Leu Val Glu Ser Met Ile Pro Arg Ser Thr Trp Lys Gly
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                725
Pro Glu Arg Leu Ala His Gly Ser Pro Phe Arg Gly Met Ser Leu Leu
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His Leu Ala Ala Gln Gly Tyr Ala Arg Leu Ile Glu Thr Leu Ser
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Gln Trp Arg Ser Val Glu Thr Gly Ser Leu Asp Leu Glu Glu Val
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                                            780
Asp Pro Leu Asn Val Asp His Phe Ser Cys Thr Pro Leu Met Trp Ala
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                                       795
Cys Ala Leu Gly His Leu Glu Ala Ala Val Leu Leu Phe Arg Trp Asn
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                                   810
Arg Gln Ala Leu Ser Ile Pro Asp Ser Leu Gly Arg Leu Pro Leu Ser
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                                825
Val Ala His Ser Arg Gly His Val Arg Leu Ala Arg Cys Leu Glu Glu
                            840
                                                845
Leu Gln Arg Gln Glu Pro Ser Val Glu Pro Pro Phe Ala Leu Ser Pro
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Pro Ser Ser Ser Pro Asp Thr Gly Leu Ser Ser Val Ser Ser Pro Ser
                                       875
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Glu Leu Ser Asp Gly Thr Phe Ser Val Thr Ser Ala Tyr Ser Ser Ala
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                                    890
Pro Asp Gly Ser Pro Pro Pro Ala Pro Leu Pro Ala Ser Glu Met Thr
                               905
Met Glu Asp Met Ala Pro Gly Gln Leu Ser Ser Gly Val Pro Glu Ala
                            920
Pro Leu Leu Met Asp Tyr Glu Ala Thr Asn Ser Lys Gly Pro Leu
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Ser Ser Leu Pro Ala Leu Pro Pro Ala Ser Asp Asp Gly Ala Ala Pro
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Glu Asp Ala Asp Ser Pro Gln Ala Val Asp Val Ile Pro Val Asp Met
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Leu Pro Phe Glu Arg Gly Arg Leu Ala Val Pro Ser Ala Pro Ser Trp
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Ala Glu Phe Leu Ser Ala Ser Thr Ser Gly Lys Met Glu Ser Asp Phe
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                            1080
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                    1110
Lys Tyr Lys Gln Leu Thr Trp Ile Ala Leu Lys Phe Ala Leu Tyr Lys
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                                                        1135
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Lys Met Thr Gln Ala Ala Ile Leu Ile Gln Ser Lys Phe Arg Ser Tyr
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            1140
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Tyr Glu Gln Lys Arg Phe Gln Gln Ser Arg Arg Ala Ala Val Leu Ile
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Gln Gln His Tyr Arg Ser Tyr Arg Arg Arg Pro Gly Pro Pro His Arg
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Thr Ser Ala Thr Leu Pro Ala Arg Asn Lys Gly Ser Phe Leu Thr Lys
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Lys Gln Asp Gln Ala Ala Arg Lys Ile Met Arg Phe Leu Arg Arg Cys
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                                                         1215
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Arg His Arg Met Arg Glu Leu Lys Gln Asn Gln Glu Leu Glu Gly Leu
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Pro Gln Pro Gly Leu Ala Thr
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Phe Pro Leu Cys Phe Leu Gly Thr Ala Phe Pro Gln Gly Glu Gln Arg
                             40
Pro Leu Glu Ala Lys Gly Leu Ala Thr Gln Gly Ala Ser Leu Pro Leu
                         55
Leu Pro Thr Val Thr Cys Val Ser Ile Lys Ser Trp Lys Met Glu Cys
Pro His Gln Gly Asp Gly Val Thr Thr Glu Ala Gly Ser Glu Leu Pro
Gln Leu Leu Gln Ala Pro Trp Pro Arg
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  720
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tcacaagaga tgagtteete agaaggeaga agaeggagae cateatetae teeegagaga
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cactcaccat aaacacatcc ccaggcagga cagatcgggg aaggggtgtg taccaggcta
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 Lys Lys Arg Phe Gln Gln Ala Thr Pro Gly Ser Ala Pro Val Ser Arg
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 Glu Gln Ala Ser Phe Leu Ala Ser Ser Phe Ser Ser Ser Ala Gly Pro
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                             40
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Leu Val Asn Arg Ile Tyr Asn Leu Gln Glu Glu Ala Arg Gln Ala Glu
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Cys Ser Thr Leu Gly Lys Asp Cys Glu Met Tyr Lys His Arg Met Asn
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Thr Val Met Leu Gln
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Ala Arg Gly Glu Gly Thr His Ser Glu Glu Glu Gly Phe Ala Met Asp
                                25
Glu Glu Asp Ser Asp Gly Glu Leu Asn Thr Trp Glu Leu Ser Glu Gly
                                                 45
Thr Asn Cys Pro Pro Lys Glu Gln Pro Gly Asp Leu Phe Asn Glu Asp
                         55
Trp Asp Ser Glu Leu Lys Ala Asp Gln Gly Asn Pro Tyr Asp Ala Asp
                                         75
Asp Ile Gln Glu Ser Ile Ser Gln Glu Leu Lys Pro Trp Val Cys Cys
                                     90
Ala Pro Gln Gly Asp Met Ile Tyr Asp Pro Ser Trp His His Pro Pro
Pro Leu Ile Pro Tyr Tyr Ser Lys Met Val Phe Glu Thr Gly Gln Phe
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                             120
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 Asp Asp Ala Glu Asp
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Ile Ser Leu Thr Met Asn Ser Lys Leu Leu Asn Gly Ser Gln Arg Val
                            40
Val Met Asp Gly Val Ile Ser Asp His Glu Cys Gln Glu Leu Gln Arg
                        55
Leu Thr Asn Val Ala Ala Thr Ser Gly Asp Gly Tyr Arg Gly Gln Thr
                                        75
65
Ser Pro His Thr Pro Asn Glu Lys Phe Tyr Gly Val Thr Val Phe Lys
Ala Leu Lys Leu Gly Gln Glu Gly Lys Val Pro Leu Gln Ser Ala His
            100
                                105
Leu Tyr Tyr Asn Val Thr Glu Lys Val Arg Arg Ile Met Glu Ser Tyr
        115
                            120
Phe Arg Leu Asp Thr Pro Leu Tyr Phe Ser Tyr Ser His Leu Val Cys
                        135
                                            140
Arg Thr Ala Ile Glu Glu Val Gln Ala Glu Arg Lys Asp Asp Ser His
                    150
                                         155
Pro Val His Val Asp Asn Cys Ile Leu Asn Ala Glu Thr Leu Val Cys
                165
                                    170
Val Lys Glu Pro Pro Ala Tyr Thr Phe Arg Asp Tyr Ser Ala Ile Leu
            180
                                185
Tyr Leu Asn Gly Asp Phe Asp Gly Gly Asn Phe Tyr Phe Thr Glu Leu
                            200
Asp Ala Lys Thr Val Thr Ala Glu Val
                        215
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gggggggag gggaagagag gggaccetgg gacceeegce eecceacce ggeegeeeet
180
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Cys Pro Glu Glu Gln Pro His Val Gly Asn Tyr Arg Leu Leu Arg Thr
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Ile Gly Lys Gly Asn Phe Ala Lys Val Lys Leu Ala Arg His Ile Leu
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Thr Gly Arg Glu Val Ala Ile Lys Ile Ile Asp Lys Thr Gln Leu Asn
                                    90
                85
Pro Ser Ser Leu Gln Lys Leu Phe Arg Glu Val Arg Ile Met Lys Gly
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Leu Asn His Pro Asn Ile Val Lys Leu Phe Glu Val Ile Glu Thr Glu
                            120
Lys Thr Leu Tyr Leu Val Met Glu Tyr Ala Ser Ala Gly Glu Pro Pro
                        135
                                             140
Thr Leu Ser Ala Leu Pro Leu Cys His Leu Pro Leu Pro Leu His Leu
                                         155
                    150
Thr Leu Thr Pro Leu Gly Leu Cys Pro Ala Gly Glu Val Phe Asp Tyr
                                    170
                165
Leu Val Ser His Gly Arg Met Lys Glu Lys Glu Ala Arg Ala Lys Phe
                                185
Arg Gln Ile Val Ser Ala Val His Tyr Cys His Gln Lys Asn Ile Val
                                                 205
                             200
His Arg Asp Leu Lys Ala Glu Asn Leu Leu Asp Ala Glu Ala Asn
                                             220
                         215
 Ile Lys Ile Ala Asp Phe Gly Phe Ser Asn Glu Phe Thr Leu Gly Ser
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 Lys Leu Asp Thr Phe Cys Gly Ser Pro Pro Tyr Ala Ala Pro Glu Leu
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 Phe Gln Gly Lys Lys Tyr Asp Gly Pro Glu Val Asp Ile Trp Ser Leu
                                 265
 Gly Val Ile Leu Tyr Thr Leu Val Ser Gly Ser Leu Pro Phe Asp Gly
                                                 285
                             280
 His Asn Leu Lys Glu Leu Arg Glu Arg Val Leu Lys Gly Lys Tyr Arg
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 Val Pro Phe Tyr Met Ser Thr Asp Cys Glu Ser Ile Leu Arg Arg Phe
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 Asp Lys Trp Ile Asn Ile Gly Tyr Glu Gly Glu Glu Leu Lys Pro Tyr
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Thr Glu Pro Glu Glu Asp Phe Gly Asp Thr Lys Arg Ile Glu Val Met
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Val Gly Met Gly Tyr Thr Arg Glu Glu Ile Lys Glu Ser Leu Thr Ser
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Gln Lys Tyr Asn Glu Val Thr Ala Thr Tyr Leu Leu Gly Arg Lys
                                        395
                    390
Thr Glu Pro Asp Glu His Gly Gly Gly Gly Ala Glu Gly Gly Ala Ala
                                    410
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Ala Arg Pro Glu Gly Glu Leu Gln His Arg Gly Glu Trp Glu Ser Arg
                                                     430
                                425
            420
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Gly Arg Asp Pro Arg Ala Ala Glu Gly Gln His Glu His Pro Arg Glu
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Ser Glu Phe Pro Ser Ser Leu Thr Gly Lys Val Ala Pro Glu Glu Phe
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Lys Gly Met Phe Ser Met Gly Trp Pro Ala Val Leu Ser Ile Thr Pro
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Thr Pro Tyr Asn Glu Asn Ile Leu Val Glu Gln Leu Tyr Met Cys Val
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Val Leu Leu Asp Leu Ala Pro Asn Leu Leu Pro Phe Gly Thr Glu
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Pro Glu Asp Lys Pro Ala Pro Lys Asn Glu Asp Glu Met Met Val Ala
Ile Phe Glu Tyr Ile Asp Arg Leu Phe Ser Ile Val Arg Pro Arg Arg
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Gln Gln Arg Ser Arg Arg Phe Arg Ala Ile Lys Glu Gly Met Glu Ala
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Thr Pro Gly Thr Glu Phe Met Asp Asn Leu Ala Lys Cys Leu Arg Tyr
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Tyr Ile Ala Asp Arg Leu Asn Asn Asp Pro Gly Trp Lys Asn Leu Thr
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Val Ile Leu Ser Asp Ala Ser Ala Pro Gly Glu Gly Glu His Lys Ile
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Met Asp Tyr Ile Arg Arg Gln Arg Ala Gln Pro Asn His Asp Pro Asn
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                                            220
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Val Lys Asp Cys Glu Gly Leu Pro Arg Glu Lys Lys Gly Lys His Asp
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Asp Gln Pro Cys Leu Cys Pro Ala Pro Ser Val Arg Thr Ala Val Ala
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Leu Thr Thr Pro Asp Ile Thr Leu Val Leu Pro Pro Asp Val Ile Gln
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Gln Glu Ala Ser Ala Leu Arg Glu Glu Thr Glu Ala Trp Ala Arg Pro
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                        455
His Glu Ser Leu Ala Arg Glu Glu Ala Leu Thr Ala Leu Gly Lys Leu
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Leu Tyr Leu Leu Asp Gly Met Leu Asp Gly Gln Val Asn Ser Gly Ile
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Gly Gln Leu Asp Arg Pro Pro Asp Leu Ala His Asp Gly Arg Ser Leu
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Trp Leu Asn Ile Arg Gly Lys Glu Ala Ala Ala Leu Ser Met Phe His
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Val Ser Thr Pro Leu Pro Val Met Thr Gly Gly Phe Leu Ser Cys Ile
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Leu Gly Leu Val Leu Pro Leu Ala Tyr Gly Phe Gln Pro Asp Leu Val
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                                                     590
Leu Val Ala Leu Gly Pro Gly His Gly Leu Gln Gly Pro His Ala Ala
Leu Leu Ala Ala Met Leu Arg Gly Leu Ala Gly Gly Arg Val Leu Ala
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Leu Leu Glu Glu Val Ser Trp Ala Gly Trp Arg Cys Cys Gly Val Gly
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Arg Gly Glu Gly Pro Val Thr Ala Ser Val Phe Ala Pro Gly Pro Glu
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Glu Arg Gly Asn Ala Ser Ser Ser Gln Lys Glu Lys Val Leu Pro Glu
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Ala Gly Ala Gly Glu Asp Ser Pro Gly Arg Asn Met Asp Thr Ala Leu
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Glu Glu Leu Gln Leu Pro Pro Asn Ala Glu Gly His Val Lys Gln Val
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                                185
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Ser Pro Arg Arg Pro Gln Ala Thr Glu Ala Val Gly Arg Pro Thr Gly
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Pro Ala Gly Leu Arg Arg Thr Pro Asn Lys Gln Pro Ser Asp His Ser
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Tyr Ala Leu Leu Asp Leu Asp Ser Leu Lys Lys Lys Leu Phe Leu Thr
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Leu Lys Glu Asn Glu Lys Leu Arg Lys Arg Leu Gln Ala Gln Arg Leu
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